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GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
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using sw model - protein search, OM protein

June 30, 2004, 19:28:48 ; Search time 17 Seconds (without alignments) 548.268 Million cell Run on:

US-10-050-552A-2 903 1 MAALQKSVSSFLMGTLATSC......EIKAIGELDLLFMSLRNACI 179 Title: Perfect score: Sequence:

BLCSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

141651 seqs, 52070155 residues Searched:

141681 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_42:* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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ALIGNMENTS

	Q9GZXb; 28-FEB-2003 (Rel. 41, Created) 28-FEB-2003 (Rel. 41, Last sequence update) 15-MAR-2004 (Rel. 43, Last annotation update) Interleukin-22 precursor (IL-22) (IL-10-related T-cell-derived	cible factor) (IL-TIF) (UNQ3099/PRO10096).	Homo sapiens (Human). Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia: Rutheria: Drimates: Catarrhini: Hominidae: Homo	196;	UENC	Van Roost E., Colau D., Renauld JC.;	rieukin-10-related T cell-derived loning and functional characteriza	stimulating factor."; Proc. Natl. Acad. Sci. U.S.A. 97:10144-10149(2000).	000	N.A. 354; PubMed=11197690; Var Boort & Colan D Amono C Wichams	C.;	"IL-TIF/IL-22: genomic organization and mapping of the human and mouse	Genes Immun. 1:488-494 (2000).	SECUENCE PROM N.A.		m o	n (IL) -22, a novel human c	<pre>interferon receptor-related proteins CRF2-4 and in-22R."; J. Biol. Chem. 275:31335-31339(2000).</pre>		SEQUENCE FROM N.A., AND VARIANT GLY-158. Rieder M.J., Carrington D.P., Chung MW., Lee K.L., Poel C.L., Yi Q.,	A. /	Submitted (JUN-2001) to the EMBL/Genbank/DLBG databases. [5]	SEQUENCE FROM N.A.	., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,	., Currell B., .Q., Hass P.E.	Kim	Watanabe C Wieand D Woods K. Xie	Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,	The secreted protein discovery initiative (SPDI), a large-scale
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10-OCT-2003 (Rel. 42, Last arnotation update)
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                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Elthe Elucopean Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by an entities requires a license agreement (See http://www.isb-resend an email to license@isb-sib.ch).
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GO:0005576; C:extracellular; IC.

GO:0005518; F:interleukin-22 receptor binding; NAS.

GO:0006953; P:acute-phase response; NAS.

GO:0006954; P:inflammatory response; NAS.

GO:0006954; P:inflammatory response; NAS.

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                                                                                                                                                                                                                                                            mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 KEASLADNNTDVRLIGEKLFRGVSAKDQCYLMKQVLNFTLEDVLLFQSDRFQFYMQEVVP
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MEDLINE=20126044; PubMed=10657629;
Dumoutier L., Louahed J., Remauld J.-C.;
"Cloning and characterization of IL-10-related T cell-derived
inducible factor (IL-TIF), a covel cytokine structurally related
IL-10 and inducible by IL-9.";
J. Immunol. 164:1814-1819(2000).
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(POTENTIAL).
(POTENTIAL).
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                                                                                                                                                                                                                                                         human
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EMBL; AJ294727; CAC19435.1; -.

NGD; MGI:1355307; I122.

GO; GO:0005515; F:protein binding; IPI.

GO; GO:0042516; P:oxygen and reactive oxygen species metabol

GO; GO:0042516; P:regulation of tyrosine phosphorylation of

InterPro; IPR000098; Interleukin_10.

SWART; SM00188; IL10; 1.

PROSITE; PS00520; INTERLEUKIN_10; 1.
                                                                                                                                                                                                                                                                                                                       inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 179;
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                                                                                                                                                                     STRAIN=129;
MEDLINE=21069354; PubMed=11197690;
Dumoutier L., Van Roost E., Colau D., Ameye G., Michaux
Renauld J.-C.;
"IL-TIE/IL-22: genomic organization and mapping of the h
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-LINKED (GLCNAC. . .) (P
ACB0ES574AA79274 CRC64;
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(Rel. 41, Last sequence update)
(Rel. 42, Last annotation update)
-22b precursor (IL-22b) (IL-10-related ctor beta) (IL-TIF beta).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
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N-LINKED (GLCNAC.
N-LINKED (GLCNAC.
                                                                                                                                                                                                                                                                                                                        to the
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larity 76.5%; Pred. No. 4.6e-61
Conservative 22; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                   SUBCELLULAR LOCATION: Secreted.
SIMILARITY: Belongs to the IL-10 family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Z
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-!- FUNCTION: Cytokine that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-FEB-2003 (Rel. 41, Creat 28-FEB-2003 (Rel. 41, Last 10-OCT-2003 (Rel. 42, Last Interleukin-22b precursor (inducible factor beta) (IL-22b OR ILTIFB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STANDARD;
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137; Conser
                                                                                                                                                      SEQUENCE FROM N.A.
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Mammalia; Eutheria;
Cervidae; Cervinae;
MCBI TaxID=9860;
                               Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                135
179 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                             SMART; SM00188; IL
PROSITE; PS00520;
                                                                                                                                                                                                                                                                                                                                                                                                                     2
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   (CSIF)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               41;
                              Bukaryota;
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DISULFID
CARBOHYD
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   factor
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T.S
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                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation—the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/armounce/or send an email to license@isb-sib.ch/.
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Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
NCBI_TaxID=10090;
                                                                            Dumoutier L., Louahed J., Renauld J.-C.,
"Cloning and characterization of IL-10-related T cell-derived inducible factor (IL-TIF), a novel cytokine structurally related IL-10 and inducible by IL-9 ".

IL-10 and inducible by IL-9 ".

J. Temunol. 164:1814-1819(2000).
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(POTENTIAE).
(POTENTIAE).
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                                                                                                                                                                                         of the human
                                                                                                                                                                                                                     inflammatory
                                                                                                                                                            .97690;
Colau D., Ameye G., Michaux L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 179;
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(IL-10) (Cytokine synthesis inhibi
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                                                                                                                                                                                         organization and mapping
                                                                                                                                                                                                                    contributes to the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                         POTENTIAL.
INTERLEUKIN-22B
                                                                                                                                                                                                                              in vivo.
SUBCELLULAR LOCATION: Secreteα.
crmiLARITY: Belongs to the IL-10 family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IL10 CEREL STANDARD; PRT; 179 AA P51746; 01-0CT-1996 (Rel. 34, Last sequence update) 01-0CT-1998 (Rel. 37, Last annotation updat I5-DEC-1998 (Rel. 37, Last annotation updat Interleukin-10 precursor (IL-10) (Cytokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ed. No. 1.16
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 714;
Pred. No.
                                                                                                                                                                                                                                                                                                                                               EMBL; AJ249492; CAB75547.1; -.
EMBL; AJ294728; CAC19436.1; -.
MGD; MGI:2151139; Iltifb.
InterPro; IPR000098; Interleukin 10.
PROSITE; PS00520; INTERLEUKIN 10; 1.
Cytokine; Glycoprotein; Signal.
                                                                    MEDLINE=20126044; PubMed=10657629;
                                                                                                                                                  STRAIN=129;
MEDLINE=21069354; PubMed=11197690
Dumoutier L., Van Roost E., Colau
Renauld J.-C.;
"IL-TIF/IL-22: genomic organizati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         20162 MW;
                                                                                                                                                                                               genes.";
Genes Immun. 1:488-494(2000)
-!- FUNCTION: Cytokine that
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179
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68
7
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54
58
97
179 Au
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136; Conser
                                                                                                                                         FROM N.A.
                                                   A.N
                                                  SEQUENCE F)
STRAIN=129
                                                                                                                                                                                                                                                                                                                                                                                                 Cytokine;
SIGNAL
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CARBOHYD
CARBOHYD
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                                                                                                                                        SEQUENCE
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Best Local
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Matches
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                           Chordata, Craniata, Vertebrata, Suteleostomi,
Cetartiodactyla, Ruminantia, Pecora, Cervoidea,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               69 NIDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEBVLFPQSDRFQPYMQEVVPFIAR-
                        Suteleostomi
                                                                                                                                                                                                                                                                    cytokines,
produced by
similarity).
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(IL-10) (Cytokine synthesis inhibitory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (POTENETAL)
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                                                                                                                                      SEQUENCE FROM N.A.

MEDLINE=96063015; PubMed=7579578;

Lockhart E., Slobbe L., Droogmans L., Griffin F., Buchan G., "The cloning and sequencing of cervine interleukin 10.";

"The cloning and sequencing of cervine interleukin 10.";

DNA Seq. 5:265-268(1995).

-!- FUNCTION: Inhibits the synthesis of a number of cytoking including IFN-gamma, IL-2, IL-3, TNF and GM-CSF produced activated macrophages and by helper T cells (By similar: "SubUNIT: Homodimer (By similarity).

-!- SUBCELLULAR LOCATION: Secreted.

-!- SIMILARITY: Belongs to the IL-10 family.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 CLLLLA-LLVQGGAAAPISSHCRLDKSNFQQPYITWRTFMLAKEASLA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; U11767; AAA85434.1; -.
HSSP; P22301; 1INR.
GO; GO:0005576; C:extracellular; NAS.
GO; GO:0005141; F:interleukin-10 receptor binding; NAS.
GO; GO:0006916; P:anti-apoptosis; ISS.
GO; GO:0006954; P:inflammatory response; ISS.
GO; GO:0006954; P:inflammatory response; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -LINKED (GLCNAC. . . . | [Pr
AlC0035D484B6050 CRC64;
                      Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 106.5; DB 1;
Pred. No. 0.0051;
2; Mismatches 74;
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(Rel. 36, Last sequence update)
(Rel. 37, Last annotation updat
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BY SIMILARITY
                        Craniata,
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Glycoprotein; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRINTS; PRO1294; INTRLEUKIN10.
ProDom; PD003687; Interleukin_10;
SMART; SM00188; IL10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Interpro; IPR000098; Interleukin_
Pfam; PF00726; IL10; 1.
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                      Chordata
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                                                                      Cervus
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P79338;
15-JUL-1998 (Rel. 36, Crea
15-JUL-1998 (Rel. 36, Last
15-DEC-1998 (Rel. 37, Last
Interleukin-10 precursor (
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Chordata; Craniata; Vertebrata; Buteleostomi
Primates; Catarrhini; Cercopithecidae;
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                                                                                                                                                                                                                                                                        GO; GO: 0005576; C:extracellular; ISS.

GO; GO: 0005576; C:extracellular; ISS.

GO; GO: 0006141; F:interleukin-10 receptor binding; ISS.

R GO; GO: 0006916; P:anti-apoptosis; ISS.

GO; GO: 0006916; P:B-cell differentiation; ISS.

GO; GO: 0042100; P:B-cell differentiation; ISS.

GO; GO: 0007253; P:Cytoplasmic sequestering of NF-kappaB; ISS.

R GO; GO: 0007253; P:cytoplasmic sequestering of NF-kappaB; ISS.

R GO; GO: 0004507; P:hemopoiesis; ISS.

R GO; GO: 004507; P:negative regulation of interferon-gamma bio.

R GO; GO: 0045130; P:negative regulation of intric cxide biosynt.

R GO; GO: 0045191; P:regulation of isotype switching; ISS.

R GO; GO: 0045092; P:T-helper 2 type immune response; ISS.

R GO; GO: 0007260; P:tyrosine phosphorylation of STAT protein; ISS.

R FIRM; PF00726; ILL0; I.

R PIRM; PF00726; ILL0; I.
                                                                                                   k/DDBJ databases.
a number of cytokines,
F and GM-CSF produced by
T cells (By similarity)
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                                                                                                     the EMBL/GenBank/DDBJ
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BY SIMILARITY.
BY SIMILARITY.
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                                                                                                            FUNCTION: Inhibits the synthesis of a nincluding IFN-gamma, IL-2, IL-3, TNF and activated macrophages and by helper T consumity: Homodimer (By similarity).

SUBCELLULAR LOCATION: Secreted.

SIMILARITY: Belongs to the IL-10 family.
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Mismatches
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Pred. No. 0.(
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Mammalia; Eutheria; Prim
Cercopithecinae; Macaca.
NCBI_TaxID=9541;
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PR01294; INTA-
: PD003687; Inter
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                                                                                           Tatsumi M.;
Submitted (FEB-1997)
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178 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genome.
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-!- SUBCELLULAR LOCATION: Secreted (Potential).
-!- SIMILARITY: Belongs to the IL-10 family.
-!- CAUTION: THERE ARE 2 BCRF1 GENES. ONE IS WRITTEN WITH A LOWER
CASE C (B(C)RF1) AND THE OTHER WITH A CAPITAL C (BCRF1).
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BCRF1, a
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ein-Barr virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Epstein-Barr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=84270667; PubMed=6087149;
Baer R., Bankier A.T., Biggin M.D., Deininger P.L., Fa Gibson T.J., Hatfull G., Hudson G.S., Satchwell S.C., Tuffnell P.S., Barrell B.G.; "DNA sequence and expression of the B95-8 Epstein-Barr Nature 310:207-211(1984).
                                                                                                                                                                                                                                                                                                                                                           virus (strain B95-8) (Human herpesvirus A viruses, no RNA stage; Herpesviridae;
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Epstein-Barr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Moore K.W., Vieira P., Fiorentino D.F., Trounstine Khan T.A., Mosmann T.R.; "Homology of cytokine synthesis inhibitory factor Epstein-Barr virus gene BCRFI."; Science 248:1230-1234(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      X-RAY CRYSTALLOGRAPHY (1.9 ANGSTROMS).
MEDLINE=97303052; PubMed=9159483;
Zdanov A., Schalk-Hihi C., Menon S., Moore K.W.,
"Crystal structure of Epstein-Barr virus protein
cellular interleukin-10.";
J. Mol. Biol. 268:460-467(1997).
                                                                                                                                                                                Æ
                                                                                                                                                                                                                            (Rel. 01, Created)
(Rel. 01, Last sequence update)
(Rel. 41, Last annotation update)
n precursor (20 kDa protein).
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SEQUENCE FROM N.A.
MEDLINE=86045995; PubMed=2998073;
Hudson G.S., Bankier A.T., Satchwell S
"The short unique region of the B95-8
Virology 147:81-98(1985).
                                                                                                                                                                                                                                                                                                                                                             Epstein-barr virus (strain B95-8) (Hum
Viruses; dsDNA viruses, no RNA stage;
Gammaherpesvirinae; Lymphocryptovirus.
NCBI_TaxID=10377;
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EMBL; M1924; AAA45900.1; -.
PIR; A03741; QQBE2.
PDB; 1VLK; 01-APR-97.
InterPro; IPR000098; Interleukin_10.
Pfam; PF00726; IL10; 1.
PRINTS; PR01294; INTRLEUKIN10.
ProDom; PD003687; Interleukin_10; 1.
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN_10; 1.
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                                                                                                                                                                                STANDARD;
                                                                                                                                                                                                                                                                                       BCRF1 protein precursor BCRF1.
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21-JUL-1986
28-FEB-2003
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P03180;
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Macaca mulatta (Rhesus macaque).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Macaca.
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produced by
similarity).
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01-OCT-1996 (Rel. 34, Last sequence update)
15-DBC-1998 (Rel. 37, Last annotation update)
Interleukin-10 precursor (IL-10) (Cytokine synthesis inhibitory
factor] {CSIP}.
                                                                                                                                                                                                                                                                                                 (POTENTIAL)
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                                                                                                                                                                                                                                                                                                 CRC64;
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MEDLINE=96003435; PubMed=7561102;
Villinger F.J., Brar S.S., Mayne A.E., Chikkala N., Ans "Comparative sequence analysis of cytokine genes from I nonhuman primates.";
J. Immunol. 155:3946-3954(1995).
-!- FUNCTION: Inhibits the synthesis of a number of cylincluding IFN-gamma, IL-2, IL-3, TNF and GM-CSF presentivated macrophages and by helper T cells (By sin activated macrophages and by helper T cells (By sin activated macrophages."
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BY SIMILARITY.
BY SIMILARITY.
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SIMILARITY: Belongs to the IL-10 family.
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bred. No. 0.
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Cytokine; Glycoprotein; Signal
                                                                                                                                                                 Pfam; PF00726; IL10; 1.
PRINTS; PR01294; INTRLEUKIN10.
ProDom; PD003687; Interleukin
SMART; SM00188; IL10; 1.
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HSSP; P22301; 1INR.
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F and GM-CSF produced by
T cells (By similarity)
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleos
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Cercocebus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=96003435; PubMed=7561102;
Villinger F.J., Brar S.S., Mayne A.E., Chikkala N., Ansari ?
"Comparative sequence analysis of cytokine genes from human nonhuman primates.";
J. Immunol. 155:3946-3954(1995).
                                                                                                                                                                                                                                                                                                                                                                7
                                                                                                                                                                                                                                                                                                                                                                                                                                                               RLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVPFI
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01-NOV-1995 (Rel. 32, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Interleukin-10 precursor (IL-10) (Cytokine synthesis
factor) (CSIF).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LSTCHIEGDDLHIQRNVQKLKDTVKKIGESGEIKAIGELDL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunol. 155:3946-3954(1995).
FUNCTION: Inhibits the synthesis of a number including IFN-gamma, IL-2, IL-3, TNF and GM-C activated macrophages and by helper T cells (SUBUNIT: Homodimer {By similarity}).
SUBCELLULAR LOCATION: Secreted.
SIMILARITY: Belongs to the IL-10 family.
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                                                                                          (GECNEC.
                           Signal, 3D-structure
POTENTIAL,
BCRF1 PROTEIN.
                                                                                                                                                                                                                                                                                                                                                            Score 103; DB Pred. No. 0.01;
                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                          N-LINKED
Glycoprotein;

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AA; 19914 MW;
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24.8%;
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CLVLLYLAPECGG-
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| P46651;
| 01-NOV-1995 (Rel
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                         Cytokine;
SIGNAL
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DISULFID
CARBOHYD
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             restrictions
                                                                                                                                                                                                                 ISS.
                                                                                     GO; GO:0005576; C:extracellular; ISS.

GO; GO:0008189; E:apoptosis inhibitor activity; ISS.

GO; GO:0006141; E:interleukin-10 receptor binding; ISS.

GO; GO:0006916; P:anti-apoptosis; ISS.

GO; GO:0006916; P:B-cell differentiation; ISS.

GO; GO:00042100; P:B-cell differentiation; ISS.

GO; GO:00042097; P:hemopoiesis; ISS.

GO; GO:0006954; P:inflammatory response; ISS.

GO; GO:0045077; P:negative regulation of interferon-gamma bio...

GO; GO:0045077; P:negative regulation of intric oxide biosynt...

GO; GO:0045191; P:negative regulation of T-cell proliferation; ISS.

GO; GO:0042092; P:T-helper 2 type immune response; ISS.

GO; GO:0007260; P:tyrosine phosphorylation of STAT protein; ISS.

InterPro; IPR000098; Interleukin_10.
                                                                                                                                                                                                                                                                                                                                                                                                                                        (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23;
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"Identification of a novel ligand-receptor pair constitutively
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              169
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            use by non-profit institutions as long as its content i modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sior send an email to license@isb-sib.ch).
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INTERLEUKIN-10.
BY SIMILARITY.
BY SIMILARITY.
N-LINKED (GLCNAC. . .) (PC
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28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Cytokine-like protein Mob-5 precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -ISSHCRLDKSN-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 101.5;
Pred. No. 0.01
0; Mismatches
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European Bioinformatics Institute.
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SEQUENCE FROM N.A.
MEDLINE=20387329; PubMed=10825166;
MEDLINE=20387329; PubMed=10825166;
                                                                                                                                                                                                                                                                                                           PF00726; IL10; 1.
3; PR01294; INTRLEUKIN10.
n; PE003687; Interleukin_10; 1.
SM00188; IL10; 1.
                                                                                                                                                                                                                                                                                                                                                                        Signal
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178 AA;
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                                                                                     P22301;
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SIGNAT
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CARBOHYD
SEQUENCE
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Best Local
Matches 4
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PRINTS;
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Isolation and expression of human cytokine synthesis inhibitory
factor cDNA clones: homology to Epstein-Barr virus open reading
BCRFI.":
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-AUG-1991 (Rel. 19, Created)
01-AUG-1991 (Rel. 19, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Interleukin-10 precursor (IL-10) (Cytokine synthesis inhibitory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE=T-cell;
WEDLINE=91142134; PubMed=1847510;
Vieira P., de Waal-Malefyt R., Dang M.-N., Johnson K.E., Kast
Vieira P., de Waal-Malefyt R., Roncarolo M.-G., Mosmann T.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      183;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CYTOKINE-LIKE PROTEIN MOB-5
N-LINKED (GLCNAC. . .) (POT)
AF7A11466C491AC4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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Submitted (OCT-1994) to the EMBL/GenBank/DDBJ databases.
[3]
SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata, Vertebrata,
Catarrhini, Hominidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11.0%; Score 99.5; DB 1
larity 27.7%; Pred. No. 0.024;
Conservative 19; Mismatches
ivated by ras oncogenes.";
siol. Chem. 275:24436-24443(2000).
SUBCELLULAR LOCATION: Secreted.
SIMILARITY: Belongs to the IL-10 family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          178 AA
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                                                                                                                                                                                                                                                                                                                                                                                                              INTERLEUKIN 10; 1.
                                                                                                                                                                                                                                                                                                           EMBL; AF269251; AAF75553.1; -...
InterPro; IPR000098; Interleukin 10.
ProDom; PD003687; Interleukin 10; 1.
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN 10; 1.
Cytokine; Glycoprotein; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NRLSTCHIEGDDLHIORNVOKLKD
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Primates;
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183
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Wammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    183 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            . Similarity
36; Conser
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SEQUENCE FROM N.A.
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Q 75
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ProDom; PD003687; Interleukin 10;
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN 10;
                    EMBL; W57627; AAA63207.1; -.

EMBL; U16720; AAA80104.1; -.

EMBL; AY029171; AAK38162.1; -.

EMBL; A38580; A38580.

PDB; IILK; 10-JUL-95.

PDB; IINK; 14-OCT-96.

PDB; IJ7V; 19-SEP-01.

PDB; ILK3; 17-JUL-02.

Genew; HGNC:5962; IL10.

MIM; 266600.
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SUBUNIT: Homodimer.

SUBCELLULAR LOCATION: Secreted.

TISSUE SPECIFICITY: Produced by a variety of cell lines, including tells, macrophages, mast cells and other cell types.

T cells, macrophages, mast cells and other cell types.

DISEASE: Defects in IL10 are a cause of susceptibility to crohn's disease (CD) [MIM:266600], a form of remitting inflammatory bowel disease (IBD). CD may involve any part of the gastrointestinal tract, but most frequently the terminal ileum arcolon. Bowel inflammation is transmural and discontinuous. Crohn'disease is commonly classified as autoimmune disease.

SIMILARITY: Belongs to the IL-10 family.
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MEDLINE=97052966; PubMed=8897595;
Zdanov A., Schalk-Hihi C., Wlodawer A.;
"Crystal structure of human interleukin-10 at 1.6-A resolution and model of a complex with its soluble receptor.";
Protein Sci. 5:1955-1962(1996).
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                                                                                                                                                                   Gesser B., Leffers H., Jinquan T., Vestergaard C., Kirstein Sindet-Pedersen S., Jensen S.L., Thestrup-Pedersen K., Larse "Identification of functional domains on human interleukin Proc. Natl. Acad. Sci. U.S.A. 94:14620-14625(1997).
                                                                                                                                                                                                                                                      DISULFIDE BONDS.
MEDLINE=93372085, PubMed=8364028,
Windsor W.T., Syto R., Tsarbopoulos A., Zhang R., Durkin J.
Windsor W.T., Syto R., Tsarbopoulos A., Zhang R., Durkin J.
Baldwin S., Paliwal S., Mui P.W., Pramanik B., Trotta P.P.;
"Disulfide bond assignments and secondary structure analysi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WARIANT CD ARG-15.
MEDLINE=22709749; PubMed=12825869;
van der Linde K., Boor P.P., Sandkuijl L.A., Meijssen M.A.,
Savelkoul H.F., Wilson J.H., de Rooij F.W.;
"A Gly15Arg mutation in the interleukin-10 gene reduces secienterleukin-10 in Crohn disease.";
interleukin-10 in Crohn disease.";
Scand. J. Gastroenterol. 38:611-617(2003).
-!- FUNCTION: Inhibits the synthesis of a number of cytokin including IFN-gamma, IL-2, IL-3, TNF and GM-CSF produced activated macrophages and by helper T cells.
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                                                                                                           the EMBL/GenBank/DDBJ databases
    CDNA and
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   human interleukin-10
                                                                             Chung M.-W.,
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MEDLINE=96173008; PubMed=8590020;
Zdanov A., Schalk-Hihi C., Gustchina A.,
               its eukaryotic expression vector.";
Harbin Yike Daxue Xue Bao 35:4-6(2001)
                                                                                                                                                                                                                                                                                                                                                                         X-RAY CRYSTALLOGRAPHY (2.0 ANGSTROMS)
MEDLINE=96032603; PubMed=7547951;
Walter M.R., Nagabhushan T.L.;
"Crystal structure of interleukin 10
                                                                                                                                                        PubMed=9405662;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biochemistry 34:12118-12125(1995)
                                                                                                                                                                                                                                                                                                                  and murine interleukin 10.";
Biochemistry 32:8807-8815(1993)
                                                                           J. P.
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(SEP-2001) to
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G -> R (in CD; decreases secretion
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GO: GO: 0005615; C: extracellular space; TAS.

GO: GO: 0005615; C: soluble fraction; TAS.

GO: GO: 0005125; F: cytokine activity; TAS.

GO: GO: 0005141; F: interleukin-10 receptor binding; NAS.

R GO: GO: 0005183; P: B-cell differentiation; NAS.

GO: GO: 0006918; P: B-cell proliferation; NAS.

R GO: GO: 0006918; P: Demotaxis; TAS.

R GO: GO: 0006919; P: Demotaxis; TAS.

R GO: GO: 0006919; P: Degative regulation of interferon-gamma bio. . .

GO: GO: 00045019; P: Degative regulation of interferon-gamma bio. . .

R GO: GO: 00045191; P: Degative regulation of cell proliferation; NAS.

GO: GO: 00045191; P: Degative regulation of cell proliferation; NAS.

GO: GO: 00045191; P: Degative regulation of cell proliferation; NAS.

GO: GO: 00045191; P: Degative regulation of cell proliferation; NAS.

GO: GO: 00045191; P: regulation of isotype switching; NAS.

GO: GO: 00045191; P: regulation of isotype switching; NAS.

GO: GO: 00045092; P: T-helper 2 type immune response; TAS.

R DitarPro: IPR000098; Interleukin_10.

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Glycoprotein; Signal; Disease mutation; 3D-structure
1 18 PCTENTIAL.
19 178 INTERLEUKIN-10.
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Interleukin-10 precursor (IL-10) (Cytckine synthesis inhibitory
factor) (CSIF).
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9; F:apoptosis inhibitor activity; ISS.
1; F:interleukin-10 receptor binding; ISS.
6; P:anti-apoptosis; ISS.
3; P:B-cell differentiation; ISS.
0; P:B-cell proliferation; ISS.
3; P:cytoplasmic sequestering of NF-kappaB; 7; P:hemopoiesis; ISS.
4; P:inflammatory response; ISS.
7; P:negative regulation of interferon-gamma? 7; P:negative regulation of MHC class II bic
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eut
Mammalia; Eutheria; Rodentia; Sciurognathi; Sciuridae;
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TISSUE=Peripheral blood;
Li D.H., Cullen J.M.;
"The woodchuck interleukin-10 gene: cloning and
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Scott B.M., O'Reilly K.L.;
Scott B.M., O'Reilly K.L.;
Submitted (OCT-1995) to the EMBL/GenBank/DDBJ databases.

-!- FUNCTION: Inhibits the synthesis of a number of cytokines, including IFN-gamma, IL-2, IL-3, TNF and GM-CSF produced by activated macrophages and by helper T cells (By similarity).

-!- SUBCELLULAR LOCATION: Secreted.

-!- SUBCELLULAR LOCATION: Secreted.
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GO; GO:0045019; P:negative regulation of nitric oxide biosynt.
GO; GO:0042130; P:negative regulation of T-cell proliferation;
GC; GO:0045191; P:regulation of isotype switching; ISS.
GC; GO:0042092; P:T-helper 2 type immune response; ISS.
GC; GO:0007260; P:tyrosine phosphorylation of STAT protein; ISS InterPro; IPR00098; Interleukin_10.
ProDom; PR01294; INTRLEUKIN10.
ProDom; PD003687; Interleukin_10; 1.
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN_10; 1.
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GO:0008189; F:apoptosis inhibitor activity; ISS.
GO:0005141; F:interleukin-10 receptor binding; ISS.
GO:0006916; P:anti-apoptosis; ISS.
GO:0030183; P:B-cell differentiation; ISS.
GO:0042100; P:B-cell proliferation; ISS.
                                                                                                                                                                                                                           CRC64;
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Mammalia; Eutheria; Carnivora; Fissipedia; Felidae;
NCBI_TaxID=9685;
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Pred. No. 0.045;
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GG: GO:0007253; P:cytoplasmic sequestering of NF-kappaB; ISS.
30; GO:0030097; P:hemopolesis; ISS.
30; GO:0006954; P:inflammatory response; ISS.
30; GO:0045347; P:negative regulation of interferon-gamma bio. . .
30; GO:0045347; P:negative regulation of mitric oxide biosynt. . .
30; GO:0045130; P:negative regulation of nitric oxide biosynt. . .
30; GO:0042130; P:negative regulation of T-cell proliferation; ISS.
30; GO:0042092; P:regulation of isotype switching; ISS.
30; GO:0042092; P:T-helper 2 type immune response; ISS.
30; GO:0007260; P:tyrosine phosphorylation of STAT protein; ISS.
EnterPro; IPR000038; Interleukin 10.
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                                                                                                                                                                                                                                                                                                                                                                                                                      01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Interleukin-10 precursor (IL-10) (Cytokine synthesis inhibit
                                                                                                                                                                                                                (POTENT
                                                                                                                                                                                                                                                      Length 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
STRAIN=PT 10000;
MEDLINE=96003435; PubMed=7561102;
Villinger F.J., Brar S.S., Mayne A.E., Chikkala N., Ansari A.
"Comparative sequence analysis of cytokine genes from human nonhuman primates.";
J. Immunol. 155:3946-3954[1995].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Butele
Mammalia; Butheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Macaca.
                                                                                                                                                                                                                                  CRC64;
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FUNCTION: Inhibits the synthesis of a number of including IFN-gamma, IL-2, IL-3, TNF and GM-CSF activated macrophages and by helper T cells (By SUBUNIT: Homodimer (By similarity).
SUBCELLULAR LOCATION: Secreted.
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N-LINKED (GLCNAC
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Pfam; PR00726; ILLO; 1.
PRINTS; PR01294; INTRLEUK
ProDom; PD003687; Interle
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178 AA;
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ROGO GO:0008189; F:apoptosis inhibitor activity; ISS.

ROGO:0008189; F:apoptosis inhibitor activity; ISS.

ROGO:0008189; F:apoptosis inhibitor activity; ISS.

ROGO:0008181; F:interleukin-10 receptor binding; ISS.

ROGO:0008183; P:B-cell differentiation; ISS.

ROGO:00082180; P:B-cell differentiation; ISS.

ROGO:00082180; P:B-cell proliferation; ISS.

ROGO:00082180; P:hemopoiesis; ISS.

ROGO:000854; P:inflammatory response; ISS.

ROGO:000854; P:inflammatory response; ISS.

ROGO:0008584; P:negative regulation of interferon-gamma bio. . .;

ROGO:00042130; P:negative regulation of nitric oxide biosynt. . .;

ROGO:00042130; P:negative regulation of T-cell proliferation; ISS.

ROGO:00042130; P:negative regulation of T-cell proliferation; ISS.

ROGO:00042092; P:T-helper 2 type immune response; ISS.

ROGO:0007260; P:tyrosine phosphorylation of STAT protein; ISS.

ROGO:0007260; ILIO; I.

RENEWRY: PROUZE; ILIO; I.

RENEWRY: PROUZE; ILIO; I.

RENEWRY: PROUZE; INTRIBUKINIO.
 (See http://www.isb-sib.ch/
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Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Boyidae; Bovinae; Bos.
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01-NOV-1995 (Rel. 32, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Interleukin-10 precursor (IL-10) (Cytokine synthesis inhibitory
factor) (CSIF).
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Pred. No. 0.056;
30; Mismatches
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INTERLEUKIN-10.
BY SIMILARITY.
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MEDLINE=94156210; PubMed=8112615;
Hash S.M., Brown W.C., Rice-Ficht A.C.;
"Characterization of a cDNA encoding bovine
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completed: June 30,
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                                                                                                                                                                                                      GO: GO: 0005576; C: extracellular; ISS.

GO: GO: 0005141; F: apoptosis inhibitor activity; ISS.

GO: GO: 0005141; F: interleukin-10 receptor binding; ISS.

GO: GO: 0005161; F: anti-apoptosis; ISS.

GO: GO: 0005161; P: anti-apoptosis; ISS.

GO: GO: 0007253; P: Cell differentiation; ISS.

GO: GO: 0007253; P: Cell proliferation; ISS.

GO: GO: 0007253; P: cytoplasmic sequestering of NF-kappaB; ISS.

GO: 0007253; P: cytoplasmic sequestering of NF-kappaB; ISS.

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GO: 00045095; P: interferon-gamma biosynthesis; IMP.

GO: 00045095; P: negative regulation of nitric oxide biosynt.

GO: 00045019; P: regulation of T-cell proliferation; ISS.

GO: 0045092; P: regulation of isotype switching; ISS.

GO: 0042092; P: T-helper 2 type, immune response; ISS.
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             LL-2, IL-3, TNF and GM-CSF pages and by helper T cells (By S: (By Similarity), ON: Secreted.
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ed. No. 0.062;
Mismatches 7
                                                                        SIMILARITY: Belongs to the IL-10 family.
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BY SIMILARITY
f expression in bovine lymphocytes.";
ene 139:257-261(1994).
!- FUNCTION: Inhibits the synthesis of
including IFN-gamma, IL-2, IL-3, TNF
activated macrophages and by helper
!- SUBUNIT: Homodimer (By similarity).
!- SUBCELLULAR LOCATION: Secreted.
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HSSP; P22301; 1INR.
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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POTENTIAL.
INTERLEUKIN-10.
BY SIMILARITY.
BY SIMILARITY.
N-LINKED (GLCNAC. . .) (POT
                                                                                                                                                                                                MEDLINE=99449065; PubMed=10520755; Wedlock D.N., Aldwell F.E., Buddle B.M.; Wedlock D.N., Aldwell F.E., Buddle B.M.; Wedlock D.N., Aldwell F.E., Buddle B.M.; Mucleotide sequence of a marsupial interleukin-10 cDNA Australian brushtail possum (Trichosurus vulpecula)."; DNA Seq. 9:239-244(1998).

-!- FUNCTION: Inhibits the synthesis of a number of cytincluding IFN-gamma, IL-2, IL-3, TNF and GM-CSF proactivated macrophages and by helper T cells (By sim activated macrophages and by helper T cells (By sim -!- SUBUNIT: Homodimer (By similarity).

-!- SUBCELLULAR LOCATION: Secreted.
    (Cytokine synthesis
                                                                Trichosurus vulpecula (Brush-tailed possum).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Metatheria; Diprotodontia; Phalangeridae;
NCBI_TaxID=9337;
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7; Mismatches 74
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HSSP; P22301; 1INR.
InterPro; IPR000098; Interleukin_10.
Pfam; PF00726; IL10; 1.
PRINTS; PR01294; INTRLEUKINIO.
ProDom; PD003687; Interleukin_10; 1.
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN_10; 1.
  (IL-10)
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1017041 segs, 315518202 residues BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table: Searched:

1017041 Total number of hits satisfying chosen parameters:

20000000000 Minimum DB seq length: Maximum DB seq length: Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

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ALIGNMENTS

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; )
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                          Last sequence update)
Last annotation update)
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Last annotation update)
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Pred. No. 0.0044;
3; Mismatches 73;
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rs I., Herbst M.M., Schountz
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21076 MW;
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Tetraodon nigroviridis (Green puffer).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
Tetradontoidea; Tetraodontidae; Tetraodon.
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SEQUENCE FROM N.A.
Lutfalla G., Roest Crollius H., Stange-Thomass.
Mogense K., Monneron D.;
"Independent expansion of a lineage-specific gene family in "Independent expansion of a lineage-specific gene family in "retebrates: The class II cytokine receptors and their ligs mammals and fish.";
Submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AY294559; AAP57417.1;
REMBL; AY294560; AAP57418.1;
"The complex of a lineage-specific gene family in submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases.

REMBL; AY294560; AAP57418.1;
"The complex of a lineage-specific gene family in light and their light and thei
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"Genomic organization of the deer mouse interleukin-10 Submitted (MAR-2003) to the EMBL/GenBank/DDBJ databases EMBL; AY251293; AAP13853.1; ".
GO; GO:0005576; C:extracellular; IEA.
GO; GO:0005125; F:cytokine activity; IEA.
GO; GO:0006955; P:immune response; IEA.
InterPro; IPR000098; Interleukin_10.
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Pred. No. 0.016;
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Ffam; PF00726; IL10; 1.
PRINTS; PR01294; INTREEUKIN10.
ProDom; PD003687; Interleukin 10;
SMART; SM00188; IL10; 1.
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SEQUENCE 178 AA; 20504 MW;
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Rao P., Jiang H., Wang F.;
"Cloning of the rhesus lymphocryptovirus viral capsid antigen and Epstein-Barr virus-encoded small RNA homologues and use in diagnosis of acute and persistent infections.";
J. Clin. Microbiol. 38:3219-3225(2000).
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MEDLINE=21602573; PubMed=11739708;
Rivailler P., Jiang H., Cho Y.-G., Quink C., Wang F.;
"Complete Nucleotide Sequence of the Rhesus Lymphocryptovirus:
Validation for an Epstein-Barr Virus Animal Model.";
J. Virol. 76:421-426(2002).
           179
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Franken M., Devergne O., Rosenzweig M., Annis B., Kieff E., "Comparative analysis identifies conserved tumor necrosis freceptor-associated factor 3 binding sites in the human and Epstein-Barr virus oncogene LMP1.";
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STRAIN=LCL8664;
MEDLINE=20304984; PubMed=10846073;
Jiang H., Cho Y.G., Wang F.;
Jiang H., Cho Y.G., Wang F.;
"Structural, functional, and genetic comparisons of E
                                                                                                                                                                                                                                                                                                                      Cercopíthicine herpesvirus 15.
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
Gammaherpesvirinae; Lymphocryptovirus.
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Last annotation update)
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MEDLINE=99412410; PubMed=10482645;
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J. Virol. 74:5921-5932(2000)
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the EMBL/GenBank/DDBJ databases
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ProDom; PD003687; Interleukin 10; 1.
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN 10; 1.
SEQUENCE 177 AA; 20545 MW; 5F940
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Submitted (JUN-1999) to
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Submitted (JAN-2000)
STRAIN=LCL8664;
Rivailler P., Quink (Submitted (MAY-1999)
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SEQUENCE FROM N.A. STRAIN=Baboon lymp

Viruses; dsDNA viruses, no RNA stage; Herpesviridae; Gammaherpesvirinae; Lymphocryptovirus. NCBI_TaxID=10394;

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STRAIN=Baboon lymphocryptovirus BA65;
Zong J.-C., Ryan J., Ling P.D., Loeb D.D.,
Hayward G.S.;
Submitted (DEC-1999) to the EMBL/GenBank/I
EMBL; AF200364; AAF23949.1; -.
HSSP; P22301; 1INR.
GO; GO:0005576; C:extracellular; IEA.
GO; GO:0006955; P:cytokine activity; IEA.
GO; GO:0006955; P:immune response; IEA.
InterPro; IPR000098; Interleukin_10.
Pfam; PF00726; IL10; 1.

Euteleostomi

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a novel cytokine-like molecule
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Tetraodon nigroviridis (Green puffer).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
Tetradontoidea; Tetraodontidae; Tetraodon.
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Lutfalla G., Roest Crollius H., Stange-Thomann N., Jaillon
Mogensen K., Monneron D.;

"Comparative genomic analysis reveals independent expansion
lineage-specific gene family in vertebrates: The class II c
receptors and their ligands in mammals and fish.";

Submitted (FEB-2003) to the EMBL/GenBank/DDBJ databases.
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Sciurognathi; Murídae;
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                                                                                      MEDLINE=21240641; PubMed=11342597;
Schaefer G., Venkataraman C., Schindler U.
"FISP (IL-4-induced secreted protein), a n
secreted by Th2 cells.";
J. Immunol. 166:5859-5863(2001).
EMBL; AF333251; AAK52470.1; ...
MGD; MGI:2135548; I124.
GO; GO:0005576; C:extracellular; IDA.
InterPro; IPR00098; Interleukin 10.
ProDom; PD003687; Interleukin 10; 1.
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN 10; 1.
SEQUENCE 220 AA; 25168 MW; SBF76C8612A
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                                                                                                                                              RTFMLAKEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPY
                                                                                                                                                                          --QLDNILLKESLJEDFKGYLGCQALSEMIQFYLEEVM-PQAEAMSLK
                                                                                                                        Gaps
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i-Poaenhded
                                                                                                                                                                                                                                                                                                                                                                                                             sostomi;
1e; Saïmiri.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                squirrel
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Cebinae
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                                                                                             Length
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                                                                                                                                                                                                                                                                                                                                                                                     Sukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Platyrrhini; Cebidae;
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Last annotation update)
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                                                                                          12;
PRINTS; PR01294; INTRLEUKIN10.
ProDom; PD003687; Interleukin_10; 1.
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN_10; 1.
SEQUENCE 171 AA; 19943 MW; 6215EDFFB259EEAA
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PRINTS; PR01294; INTRLEUKIN10.
ProDom; PD003687; Interleukin 10; 1.
SMART; SM00188; ILIO; 1.
PROSITE; PS00520; INTERLEUKIN 10; 1.
SEQUENCE 178 AA; 20432 MW; E65087383C0FD6FA
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                                                                                        9.5; DB 1
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26.9%; Pred. No. 0.086;
ive 21; Mismatches
                                                                                     11.0%; Score 99.5; I
llarity 27.7%; Pred. No. 0.06
Conservative 20; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
MEDLINE=21972723; PubMed=11976788;
Heraud J.M., Lavergne A., Kazanji M.;
"Molecular cloning, characterization, and
monkey (Saimiri sciureus) Th1 and Th2 cytc
Immunogenetics 54:20-29(2002).
EMBL; AF294758; AAK92045.1; -.
GO; GO:0005576; C:extracellular; IEA.
GO; GO:0005125; F:cytokine activity; IEA.
GO; GO:0005125; F:cytokine activity; IEA.
InterPro; IPR000098; Interleukin_10.
Pfam; PF00726; IL10; 1.
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33; Conser
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Best Local Similarity
Matches 32; Conser
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01-OCT-2002 (
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Crollius H.,

Lutfalla G., Roest Croll Mogense K., Monneron D.;

Mogense K., N "Independent

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                                                                                                                                                                                                                                                                                                                                                                                                  -FPQSDRFQPYMQEVVPFIARISNR
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                                                                                                                                                                                                                                                                                                                                                         LESVILLIESSSCAVWCATLCNNRCCSFLEGFPARLKVLRENYSNIREYYEANDDLDLVLI
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                                                                                                                                                                                                                                                                                                     Gaps
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    and their ligands
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                                                                                                                                                                                                                                                                  Length 184;
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                                                                                                                                                                                                                                                                                                   Indels
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                                  the EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cyprinus carpio (Common carp).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eu
Actinopterygii; Neopterygii; Teleostei; Ostariophysi;
Cyprinidae; Cyprinus.
NCBI_TaxID=796?.
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                                                                                                                                                                                                                                                                                                87.
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Last annotation updat
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                                                                                                                                                                                                                                                            Similarity 21.3%; Score 97; DB 13; Similarity 21.3%; Pred. No. 0.13; 5; Conservative 34; Mismatches 81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7FE85116831FDB39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13;
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Pred. No. 0.16;
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                                          EMBL; AJ544312,
EMBL; AJ544898; CAD67786.1; -.
EMBL; AY294557; AAP57415.1; -.
GO; GO:0005576; C:extracellular; IEA.
GO; GO:0006955; P:immune response; IEA.
InterPro; IPR00098; Interleukin_10.
Pfam; PF00726; IL10; 1.
PRINTS; PRC1294; INTRLEUKIN10.
ProDom; PD003687; Interleukin_10; 1.
ProDom; PD003687; Interleukin_10; 1.
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                                 2003) to the CAD67773.1; CAD67786.1;
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larity 24.3%;
Conservative 2
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                                  Submitted (MAY-2003)
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28; Conser
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              fish
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01-OCT-2003
01-OCT-2003
01-OCT-2003
                 mammals and
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Best Local S
Matches 35
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--ISNRLSTCHIEGDDLHIORN
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                                                                                                                                                                                                                                               Pehr
                                                                                                                                  Buteleostomi;
Felis.
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                                                                                                                                                                                                       SEQUENCE FROM N.A.

Leutenegger C.M., Huder J.B., Mislin C., Hofmann R., Fe
Leutenegger C.M., Huder J.B., Mislin C., Hofmann R., Fe
Dean G.A., Higgins J., Pedersen N.C., Lutz H.;

"Molecular cloning and expression of feline interleukin-10.";

"Molecular cloning and expression interleukin-10.";

"Molecular cloning and expression interleukin-10.";

"Molecular cloning and expression interleukin-10.";

Resident Molecular cloning interleukin-10;

"Probom; PD003687; Interleukin-10; 1.

SMART; SM00188; IL10; 1.

PROSITE; PS00520; INTERLEUKIN-10; 1.

SEQUENCE 178 AA; 20575 MW; 86A3FE97D9C18D08 CRC64;
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Madireddi M.T., Lin J., Su Z., Shay J.W., Huberman E., Fisher Madireddi M.T., Lin J., Su Z., Shay J.W., Huberman E., Fisher Tenomic Structure, Chromosomal Localization and Expression of Melanoma Differentiation Associated Gene-7 (mda-7): Potential Relationship with Cellular Senescence.";
Submitted (FEB-2000) to the EMBL/GenBank/DDBJ databases.

EMBL; AF235006; AAK52590.1; -..

MGD; MGI:2135548; Il24.

GO:0005576; C:extracellular; IDA.
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Sciurognathi, Muridae, Murinae,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              178;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
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05CA43872D53D555 CRC64;
                                                                                   IL-10.
Felis silvestris catus (Cat).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Carnivora; Fissipedia; Felidae;
NCBI_TaxID=9685;
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 95.5; DB Pred. No. 0.17;
    Created)
Last sequence up
Last annotation
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01-DEC-2001 (TrEMBLrel. 19, Last sequence of the collocation of the collocation (TrEMBLrel. 25, Last annotation Melanoma differentiation associated gene-7
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ProDom; PD003687; Interleukin 10; 1
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN 10; 1.
SEQUENCE 181 AA; 20812 MW; 05CAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              145 VOKLKDTVKKIGESGEIKAIGELDL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score
Pred.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10.6%;
llarity 29.4%;
Conservative 2
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13,
24,
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25; Conser
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Best Loca≧ Similarity
                                                               Interleukin-10.
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01-MAY-2000
01-MAY-2000
01-JUN-2003
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SEQUENCE
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  TaxID=9986;
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                                                                                                          ISNRLSTCHIE
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                                   --VOGGAAAPI-SSHCRLDKSNFQQPYITNRTFMLAKEASLADNNTDVRLI
                                                                     -TGVVLPELWEAFWTVKNTVQTQDDITSIRLL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       --KVL
   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                               170
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-FKNYHSKIAKF--KVLRSFST-
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O9TSJ4;
O9TSJ4;
O9TSJ4;
O1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
O1-MAY-2003 (TrEMBLrel. 24, Last annotation update)
Interleukin-10.
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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GEVDIL
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 37;
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                                                                                                      GEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVPFIAR
                                                                                                                                                                                                            117 NNFIVIMSQLQPSKDNSMLPISESAHQRFLLFRRAFKQLDTEVALVKAF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Longaker M.T., Bertolami
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Last annotation update)
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associated gene, m
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68
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171
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Pred. No. 0.18
9; Mismatches
  Mismarches
                                                                                                                                                                                                                                                                                                       183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GO; GO: 0005576; C: extracellular; IEA.
GO; GO: 0005125; F: cytokine activity; II
GO; GO: 0006955; P: immune response; IEA
InterPro; IPR000098; Interleukin 10.
ProDom; PD003687; Interleukin 10; 1.
PROSITE; PS00520; INTERLEUKIN 10; 1.
                                                                   10 CLSLILLLWNQVPGLEGQEFRSGSCQV-
                                                                                                                                      68 KPQVLRNVSGAESCYLAHSLLKFYLNTV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=Sprague-Dawley;
MEDLINE=99308986; PubMed=10381256;
Soo C., Shaw M.M., Freymiller E., LocChiu R., Tieu A., Ting K.;
"Cutaneous rat wounds express c49a, human melanoma differentiation assoc J. Cell. Biochem. 74:1-10(1999).
EMBL; AF004774; AAB69171.1;
                                                                                                                                                                           GDDLHIQRNVQKLKD-TVKKIGESGE-
                                                                                                                                                                                                                                                                                                                                          Created
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(3)
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10116;
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larity 26.9%;
Conservative 1
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24.
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                                 20 CLLLLALL-
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35; Conser
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Q9WVP8;
01-NOV-1999
01-NOV-1999
 42;
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14
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Matches
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Herbst M.M., Schountz T.;

Herbst M.M., Schountz T.;

Herbst M.M., Schountz T.;

"Cloning of the deer mouse interferon gamma, interleukin-10 and tumminerosis factor genes.";

Endited (SEP-2000) to the EMBL/GenBank/DDBJ databases.

R BMBL; AF307012; AAG30263.1; -.

R HSSP; P22301; IINR.

GO; GC:0005576; C:extracellular; IEA.

GO; GC:0005125; F:cytokine activity; IEA.

GO; GO:0006955; P:immune response; IEA.

GO; GO:0006955; P:immune response; IEA.

R GO; GO:0006955; P:immune response; IEA.

R GO; GO:000698; Interleukin_10.

R PRINTS; PR01294; INTRLEUKIN10.

R PRINTS; PR01294; Interleukin_10; 1.

SMART; SM00188; IL10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 SSALLCCLVFLGGTGASRGODTPAENSCIHFPGGLPHMLRELRAAFGRVKTFFQSKD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Euteleostomi; Sigmodontinae
                                                      P.J.;
and IL-10 from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 178;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ---ISNRLSTCHIEGDDLHIQRNVQKLKDTVKKIGESGEIKAIGELDL
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                                                                                                                                                                                                                                                                     InterPro; IPR000098; Interleukin_10.
Pfam; PF00726; IL10; 1.
PRINTS; PR01294; INTRLEUKIN10.
ProDom; PD003687; Interleukin_10; 1.
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN_10; 1.
SEQUENCE 178 AA; 20146 MW; A5E0B2A8F04D5170 CRC64;
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Interleukin-10 (Fragment).
IL10.
Peromyscus maniculatus (Deer mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
                                                         Kerr
IL-6
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Pred. No. 0.24;
10; Mismatches
                                             Perkins H.D., van Leeuwen B.H., Hardy C.M., "The complete cDNA sequences of IL-2, IL-4, European rabbit (Oryctolagus cuniculus)."; Cytokine 12:555-565(2000).

EMBL; AF068058; AAC23839:1; -.
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                                                                                                                                                                                                                          IBA.
                                                                                                                                                                      HSSP; P22301; IINR.
GO; GO:0005576; C:extracellular; IEA.
GO; GO:0005125; F:cytokine activity; II
GO; GO:0006955; P:immune response; IEA.
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TISSUE=Lymph node;
MEDLINE=20304414; PubMed=10843729;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN_10;
NON_TER 1 1
NON_TER 131
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Best Local Similarity 22.0%;
Matches 37; Conservative 30
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131 AA; 15398 MW; GEAGEA20CDC0234F CRC64;
                              Query Match
Best Local Similarity
SQ SEQUENCE
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	114	77	<u>ი</u>	30	
\$ 6 0 0	FPQSDRFQPY	: ENHGPE	GELDL 1	SEFDI 1	
131;	VLFPQS	: : VM-PQA	GEIKAI	 GWIKAM	
10.2%; Score 92.5; DB 11; Length 131; 24.4%; Pred. No. 0.23; ive 25; Mismatches 54, Indele 11	VLNFTLEE VLNFTLEE	19 KTFFQKKDQLDSILLTDSLMKDFKGYLGCQALSEMIQFYLVEVM-PQAENHGPE 71	115 MQEVVPFIARISNRLSTCHIEGDDLHIQRNVQKLKDTVKKIGESGEIKAIGELDL 169		
: H	YEMKO	: : QALSEI	OKLKD.	:::{ EQVKS1	
, DB 0.23; hee	MSERÇ	GXLGC	IQRNV	: KSKAV	
92.5 No. 5	FHGVS	MKDFK	СББЕН	LPCEN	,
Score Pred.	IGEK	LTDSL	TCHIE	RCHRE	Ċ
2.4 %%;	TDVRI	 	SMRLS	RRRIC	•
10. 24.	LADN)	1	KLKTI	ć
10.2%; Score 92.5; DB larity 24.4%; Pred. No. 0.23; Conservative 25; Mismatches	MLAKEAS	FOKED	VVPFIAE	HLNFLGE	(;
100 S	. S. R	S KTE	S MOE	72 EKB	4
Query Match Best Focal Matches 22		1	1.1	1~	[4000
Query X Best Eo Matches	λo	DP	Š	D.D.	Sparch completely. True of order

Search completed: June 30, 2004, 19:36:04 Job time: 43 secs

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GenCore (c) 1993
            Copyright
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model ¥, W using search, - protein OM protein

23 Seconds June 35, 2004, 19:27:38 Run on:

Search time 23 Second (without alignments) 401.785 Million cell

updates/sec

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MAALOKSVSSFLMGTLATSC. US-10-050-552A-2 903 score: Title: Perfect so Sequence:

BLOSUM62 Gapop 10.0 , Scoring table:

residues 389414 segs, 51625971 Searched:

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Gapext

389414 satisfying chosen parameters: hits O H number Total

2000000000 seg length: seg length: Minimum Maximum

100% 45 g Post-processing: Minimum Match 04 Maximum Match 104 Listing first 45

summaries

Database

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	Description	Sequence 2, Appl	-28 Sequence 28, App	-28 Sequence 28, App	B-15 Sequence 15, App.	-27 Sequence 27, App	-27 Sequence 27, App	-16 Sequence 16, App	Sequence 2, Appl	-2 Sequence 2, Appl	Sequence 2, Appl	2, Appl	Sequence 4, Appl	2, Appl	26 Sequence 26, App		Sequence 2, Appl	2, Appl	Sequence 26, App	Sequence 3, Appl	2, Appl	-1 Sequence 1, Appl	Seguence 1,	Sequence 1,	Sequence 1, Appl	equence 6,	Sequence 1, Appl	
SUMMARIES	E	S-09-870-574	19-568	-354-243	-09-178-973	-419-568	-09-354-243	-09-178-973	-08-270-805	-08 - 410 - 654	-08-474-851	-08-481-560	-08-934-959	-08-170-113	-08-765-094	2-797	-08-643-810	-09-552-613	-09-512-256	-09-452-624	8-884-077	-08-270-805	-08-410-654	-851	-08-481-560	-08-934-959		400 HUA 00
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US-09-082-797-25	-643-	US-09-552-613-1	-512-2	US-09-462-941-16	US-08-270-805C-4	US-08-410-654B-4	US-08-474-851-4		-170-1	US-08-643-810A-4	US-09-552-613-4	PCT-US93-07646-2	-09-452-624A-	89	4	H	US-08-481-560-3
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ENCODING PANCREATIC DISORDER
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US-09-870-574-2
| Sequence 2, Application US/09870574
| Patent No. 6551799
| Patent No. 6551799
| GENBRAL INFORMATION:
| APPLICANT: Aggarwal, Sudeepta APPLICANT: Aggarwal, Sudeepta APPLICANT: Aggarwal, Sudeepta APPLICANT: Aggarwal, Sudeepta APPLICANT: Foster, Jessica S.
| APPLICANT: Goddard, Audrey
| APPLICANT: Goddard, Audrey
| APPLICANT: Foster, Jessica S.
| APPLICANT:
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Pred. No. 2.2e-100;
3; Mismatches 0;
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Best Local Similarity
Matches 176; Conser
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LENGTH: 179
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RESULT 2 US-09-419-568F-28 ; Sequence 28, Application US/09419568F

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APPLICANT: Dumoutier, Laure
APPLICANT: Loubed, Jamila
APPLICANT: Renauld, Jean-Christophe
TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode
TITLE OF INVENTION: (TIPS)
TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
FILE REFERENCE: LUD 5543
CURRENT APPLICATION NUMBER: US/09/178,973B
CURRENT FILING DATE: 1998-10-26
NUMBER OF SEQ ID NOS: 17
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Pred, No. 9.5e-79;
22; Mismatches 20;
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Pred. No. 9.5e
2; Mismatches
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APPLICANT: Dumoutier, Laure
APPLICANT: Louhed, Jamila
APPLICANT: Louhed, Jean-Christophe
TITLE OF INVENTION: Isolated Nucleic Acid Mol
TITLE OF INVENTION: (TIFS) The Proteins Ence
FILE REFERENCE: LUD 5543.2
CURRENT APPLICATION NUMBER: US/09/419,568F
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Patent No. 6331613
                                                                                                             US-09-178-973H 15
# Sequence 15 Application US/09178973|
; Patent No: 6274710
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137; ConsetVative
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PRIOR FILING DATE: 1999-0
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PRIOR FILING DATE: 1998-1
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FNGTH: 179
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                                                                          Cell
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                         APPLICANT: Dumoutier, Laure
APPLICANT: Bounoutier, Laure
APPLICANT: Renauld, Jean-Christophe
TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode
FILE REFERENCE: LUD 5543.2
CURRENT APPLICATION NUMBER: US/09/419,568F
CURRENT FILING DATE: 1999-10-18
PRICR APPLICATION NUMBER: US09/354,243
PRICR PILING DATE: 1999-07-16
PRICR PILING DATE: 1999-07-16
PRICR PILING DATE: 1999-10-26
NUMBER OF SEQ ID NOS: 29
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; Patent No. 6359117
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Louhed, Jamila
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: (TIFS)
; TITLE OF INVENTION: (TIFS)
; TITLE OF INVENTION: (TIFS)
; TITLE OF INVENTION: The Proteins Encoded, and Uses Ther
; FILE REFERENCE: LUD 5543.1
; CURRENT APPLICATION NUMBER: US/09/354,243B
; CURRENT FILING DATE: 1999-07-16
; PRIOR FILING DATE: 1999-10-26
                                                                                                                                                                                                                                                                                                                                 Score 875; DB 4; I
Pred. No. 9.9e-98;
; Mismatches 2;
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Pred. No. 9.9e-98;
; Mismatches 2;
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ilarity 96.1%;
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                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 172; Conser
 Patent No. 6331613
GENERAL INFORMATION:
6331613
                                                                                                                                                                                                                                                                                            ; FEATORE:
JS-09-419-568F-28
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354-243B-28
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LENGTH: 179
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LENGTH: 179
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KEASLADNNTDVRLIGEKLFRGVSAKDOCYLMKOVLNFTLEDILLPOSDRFRPYMQEVVP
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                                                                                               1 MAALOKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNPQQPYITNRTFMLA
                                                                                                                Gaps
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68 :
                                         DB 3;
2.9e-78;
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Pred. No. 0.0002;
7; Mismatches 6
                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: Macintosh
COMPUTER: Macintosh
OPERATING SYSTEM: 7.5.3
CURRENT APPLICATION DETA:
APPLICATION NUMBER: US/08/270,805C
FILING DATE: 05-July-1994
CLASSIFICATION: 424
                                        Score 714;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/995,564
FILING DATE: 23-Dec-1992
APPLICATION NUMBER: US 07/830,493
FILING DATE: 04-Feb-1992
APPLICATION NUMBER: US 07/641,342
FILING DATE: 16-Jan-1991
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/08270805C
Patent No. 5776451
GENERAL INFORMATION:
APPLICANT: Di-Hwei Hsu
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Schering-Plough C
STREET: 2000 Galloping Hill
CITY: Kenilworth
STATE: New Jersey
                                                                   23
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TELEPHONE: 908-298-2987
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illarity 24.8%;
Conservative 2
                                       79.18;
76.08;
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                                                                                                                                                                                                                                                                                                                                                                              Moore
                                                                     Conservative
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TELEFAX: 908-298-5388
INFORMATION FOR SEQ ID NO:
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CRGANISM: Mus musculus US-09-178-973B-16
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MOLECULE TYPE:
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APPLICANT: Renauld, Jean-Christophe
TITLE OF INVENTION: [TIF8]
TITLE OF INVENTION: (TIF8)
TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
FILE REFERENCE: LUD 5543
CURRENT APPLICATION NUMBER: US/09/178,973B
CURRENT FILING DATE: 1998-10-26
                                                                                                                                                                                                                                                                                        Louhed, Jamila
Renauld, Jean-Christophe
FENTION: Isolated Nucleic Acid Molecules which En
                                                                    KEASILADINITDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDR.
                                                                                                                             KERSLADINI DVRLIGEKLFHGVSMSERCYLMKOVLNETLEEVLFPOSDR
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Best Local Similarity, O 76.5%; Pred. No. 9.5e-79
Matches 137; Conservative 22; Mismatches 2
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                                                                                                                                                                                                                            Sequence 27, Application US/29354243B
Patent No. (150117)
GENERAL INFORMATION:
APPLICANT: Dumoutier, Laure
APPLICANT: Loubed, Jamila
APPLICANT: Renauld, Jean-Christophe
TITLE OF INVENTION: Isolated Nucleic Acid Mo
TITLE OF INVENTION: (TIFs)
TITLE OF INVENTION: (TIFs)
TITLE OF INVENTION: The Proteins Encoded, an FILE REFERENCE: LUD 5543.1
CURRENT APPLICATION NUMBER: US/09/354,243B
PRIOR APPLICATION NUMBER: US09/178,973
PRIOR FILING DATE: 1998-10-26
NUMBER OF CENTIN NO. 10 NO.
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Patent No. 6274710
GENERAL INFORMATION:
APPLICANT: Dumoutier, Laure
APPLICANT: Louhed, Jamila
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US-09-178-973B-16
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US-09-354-243B-27
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US-09-354-243B-27
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LENGTH: 179
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LENGTH: 179
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                                                                                                                                                                                                                                                                    of An Interleukin-10 Antagonist t
Cell Mediated Autoimmune Disorder
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                                               LSTCHIEGDDLHIQRNVQKLKDTVKKIGESGEIKAIGELDE
                                                                 LRRCHRFLPCENKSKAVEQIKNAFNKLQEKGIYKAMSEFDI
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Pred. No. 0.00
7; Mismatches
                                                                                                                                                                                                                                                                                                                             SEE: Schering-Plough Corporation
: 2000 Galloping Hill Road
Kenilworth
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MRBER: US 08/229,854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08/229,854 FILING DATE: 19-APR-1994 APPLICATION NUMBER: US 07/926,853 FILING DATE: 06-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'A:
US/08/474,851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 06-AUG-1992
APPLICATION NUMBER: US 07/742,12:
FILING DATE: 06-AUG-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                     APPLICANT: Rene de Waal Malefyt
APPLICANT: Di-Hwei Hau
APPLICANT: Anne O'Garra
APPLICANT: Hergen Spits
TITLE OF INVENTION: USE Of An Int
TITLE OF INVENTION: A B Cell Medi
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e 2, Application US/08474851
No. 5837232
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Microsoft Word 6.0
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TELEPHONE: 908-298-2987
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 07-JUN-199
CLASSIFICATION: 424
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SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
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OPERATING SYSTEM: 7
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FILING DATE: 24-MA
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                                                                                                                                                           Sequence 2, Application
Patent No. 5837232
GENERAL INFORMATION:
APPLICANT: Rene de
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US-08-474-851-2
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Matches 40
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       -PYITNRTFMLAKEASLADMNIDV
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                                                                                             57 LLLKESLLEDFKGYLGCQALSEMIQFYLEEVM-PQAENQDPEAKDHVMSL
                                    -TDQC----DNFPQMLRDLRDAFSRVKTFFQTKI
                                                                    RLIGEKLFHGVSMSERCYLMKQVLNFTLSEVLFPQSDRFQPYMQEVVPF3
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                                                                                                                                                                                                                                    Sequence 2, Application US/08410654B

Patent No. 5833976

GENERAL INFORMATION:

APPLICANT: Rene de Waal Malefyt

APPLICANT: Di-Hwei Hsu

APPLICANT: Anne O'Garra

APPLICANT: Hergen Spits

TITLE OF INVENTION: Septic Shock

NUMBER OF SEQUENCES: 61

CORRESPONDENCE ADDRESS:
                                                                                                                                129 LSTCHIEGDDIHIORNVQKLKDTVKKIGESGEIKAIGELDL 169
                                                                                                                                                Score 103; DB 2
Pred. No. 0.0002
7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Schering-Plough Corporation
STREET: 2000 Galloping Hill Road
CITY: Kenilworth
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20 CLLLLALLVQGGAAAPISSHCRLDKSNFQQ-
        CLLLLALLVQGGAAAPISSHCRLDKSNFQQ
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APPLICATION NUMBER: US/08/410,654B
FILING DATE: 24-MAR-1995
CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/229,854

RILING DATE: 19-APR-1994
APPLICATION NUMBER: US 07/926,853
FILING DATE: 06-AUG-1992
APPLICATION NUMBER: US 07/742,129
FILING DATE: 06-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: FOULKe, Cyrichia L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MBER: US/08/410,654B
24-MAR-1995
                                                                                                                                                                                                          7 9
1-410-654B-2
nence 2, Application US/08410654B
-ent No. 5833976
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Microsoft Word 5.1a
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TELEPHONE: 908-298-2987
TELEFAX: 908-298-5388
FORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: 7.5.3
SOFTWARE: Microsoft Word
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Foulke, Cyrchia L. REGISTRATION NUMBER: 32, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 170 amino acids amino acids
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CLVLLYLAPECGG-
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INFORMATION FOR
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Best Local S
Matches 40
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US-08-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 170;
                                                                                                                      APPLICANT: Knappe, Andrea
APPLICANT: Fickenscher, Helmut
APPLICANT: Fleckenstein, Bernhard
TITLE OF INVENTION: MAMMALIAN CYTOKINE; RELATED REAGENTS
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
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156
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                                                                                                                                                                                                                                                                                                                                                                                                Version #1.30
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Pred. No. 0.0002;
; Mismatches 6
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                                                                                                                                                                                                                               DNAX Research Institute
L'California Avenue
                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/934,959
FILING DATE: 22-SEP-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MMBER: US 60/027,368
23-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DX0644K
                                                                                                                                                                                                                                                                                                                                                                                           Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                 IBM PC compatible YSTEM: PC-DOS/MS-DOS
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Moore, Kevin W.
de Waal Malefyt, Rene
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Patent No. 6106823
GENERAL INFORMATION:
APPLICANT: Vieira, Paulo J.
APPLICANT: Moore, Kevin W.
APPLICANT: de Waal Malefyt, Ren
                                                                                   US/08934959
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Ching, Edwin P.
REGISTRATION NUMBER: 34,090
REFERENCE/DOCKET NUMBER: D)
TELECOMMUNICATION INFORMATION
TELEPHONE: 650-852-9196
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| Similarity 24.8%;
40; Conservative 2
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11 CLVLLYLAPECGG--
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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                                                                             4, Application
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California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
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OPERATING SYSTEM:
SOFTWARE: Patent
                                                                                  Sequence 4, Application Patent No. 5989867 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                    USA
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US-08-934-959-4
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                                                                                                                                                                                                                                   ADDRESSEE:
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CITY: Pa
STATE: C
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                                                   RESULT 12
US-08-934-959-4
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                                                                                                                                                                                                                    APPLICANT: Hergen Spits
TITLE OF INVENTION: Use of Interleukin-10 to Modulate
TITLE OF INVENTION: Inflammation or T-Cell Mediated
TITLE OF INVENTION: Immune Function
                          129 LSTCHIEGDDLHIQRNVQKLKDTVKKIGESGEIKAIGELDL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        68;
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Pred. No. 0.0002
7; Mismatches
                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Schering-Plough Corporation
STREET: 2000 Galloping Hill Road
CITY: Kenilworth
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 CLLLLALIVQGGAAAPISSHCRLDKSNFQQ-
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PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/410,654

FILING DATE: 24-MAR-1995

APPLICATION NUMBER: US 08/229,854

APPLICATION NUMBER: US 07/926,853

FILING DATE: 06-AUG-1992

APPLICATION NUMBER: US 07/742,129

FILING DATE: 06-AUG-1991

ATTORNEY/AGENT INFORMATION:

NAME: 06-AUG-1991
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FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
                                                                                                                       Sequence 2, Application US/08481560 Patent No. 5837293 GENERAL INFORMATION:
                                                                                                                                                                        Malefyt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: 7.5.3
SOFTWARE: Microsoft Word 6.0
CURRENT APPLICATION DATA:
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TELEPHONE: 908-298-2987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Foulke, Cynthia L. REGISTRATION NUMBER: 32,364 REFERENCE/DOCKET NUMBER: DX
                                                                                                                                                                                                                                                                                                                                                                                      COUNTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
TITEL Macintosh
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ilarity 24.8%;
Conservative 27
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amino acid
                                                                                                                                                                     Rene de Waal
Di-Hwei Hsu
Anne O'Garra
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R SEO ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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INFORMATION FOR
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Patent No. 6159937
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLVLLYLAPECGG----TDQC----DNFPQMLRDLRDAFSRVKTFFQTKD-
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J
D
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               Fluckinger, Anne-Catherine
Banchereau, Jacques
VENTION: TREATMENT OF NEOPLASTIC DISEASE WITH
VENTION: INTERLEUKIN-10
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                                                                                                                                                                                                                                                                              Version #1.25
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Pred. No. 0.0002;
7; Mismatches 66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 04-FEB-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/641,347
FILING DATE: 16-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Ching Paris
                                                                                                                                                                                                                                                                                                                      FILING DATE: 17-DEC-1993
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/933,419
FILING DATE: 21-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/020,018
FILING DATE: 12-JUL-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/020,018
FILING DATE: 17-FEB-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/020,018
FILING DATE: 04-FEB-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Ching, Edwin P.
REGISTRATION NUMBER: 34,090
REFERENCE/DOCKET NUMBER: DX0116Q1KX
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-852-9196
                                                                                                              DNAX Research Institute
. California Avenue
                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/170,113
FILING DATE: 17-DEC-1993
                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27;
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Conservative 27
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                        APPLICAMT: Banchereau,
TITLE OF INVENTION: TRE
TITLE OF INVENTION: INT
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
de Vries,
                                                                                                                                            Palo Alto
California
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; MOLECULE TYPE:
US-08-170-113-2
                                                                                                               ADDRESSEE:
                                                                                                                                                                            COUNTRY:
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               PPLICANT:
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STATE:
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Pred. No. 0.0002;
7; Mismatches 66
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                                                                           P.L.L.C.
, Suite 300
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R: GRONHOJ-LARSEN=1
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                                                                                                                                                                                                                                                                                                 FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION APPLICATION
  Christian
                               IMMUNOMODULATORS 2.7
                                                                                                                                                                                                                                                            MBER: US/08/765,094C
06-JAN-1997
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CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK, P.
STREET: 419 Seventh Street N.W.,
                                                                          SROWDY AND NEIMARK, P
Seventh Street N.W.,
                                                                                                                                                                                               COMPUTER: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 26, Application US/09082797
Patent No. 6168791
GENERAL INFORMATION:
APPLICANT: GRONHOJ LARSEN, Chris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27;
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illarity 24.8%;
Conservative 2
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TELEFAX: (202)737-3528
INFORMATION FOR SEQ ID NO: 20
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 05-JUL-1994 ATTORNEY/AGENT INFORMATION:
  LARSEN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GRONHOJ LARSEN,
GESSER, Borbala
                                                                                                                                                                                                                                                                                                                                                                                                  NAME: COOPER, IVET P.
REGISTRATION NUMBER: 28,
REFERENCE/DOCKET NUMBER:
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nino acid
                 Borbal
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                             TITLE OF INVENTION: IN NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                            APPLICATION NUMBER:
FILING DATE: 06-JA
                                                                            BROWDY
GESSER,
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Washington
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TOPOLOGY: linear
MOLECULE TYPE: pep
                                                                                                        Washington
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Best Local Similarity
Matches 40; Conser
                                                                                                                                     U.S.A.
                                                                                          419
                                                                                                                                COUNTRY: U.
                                                                          ADDRESSEE:
STREET: 41
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APPLICANT:
APPLICANT:
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Sequence 26, Application US/08765094C

RESULT 14 US-08-765-094C-26

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--PYITHRIFMLAKEASLADNNTDV 72
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                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/082,797
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 24.8%; Fred. No. 0.0002;
Matches 40; Conservative 27; Mismatches 68
                                                                                                                                                                                                                                                                                                                                                 28,005
R: GRONHOJ-LARSEN=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 CLILLALLVQGGAAAPISSHCRLDKSNFQQ-
                                                                                                                                                                               FILING DATE:

PRIOR APPLICATION NUMBER: 08/765,094
FILING DATE:

PRIOR APPLICATION NUMBER: 08/765,094
FILING DATE:

APPLICATION NUMBER: DK 0800/94
FILING DATE: 05-JUL-1994
ATTORNEY/AGENT INFORMATION:

NAME: COOPER, IVER P.

REGISTRATION NUMBER: 28,005
REFERENCE/DOCKET NUMBER: GRONHOJ-
TELEPHONE: (202) 628-5197
TELEPHONE: (202) 737-3528
INFORMATION FOR SEQ ID NO: 26:

SEQUENCE CHARACTERISTICS:
LENGTH: 170 amino acids

TYPE: amino acids

TYPE: amino acids

TYPE: amino acids

TOPOLOGY: linear

MOLECULE TYPE: peptide

US-09-082-797-26
              D.C.
: U.S.A.
STATE: D.C. COUNTRY: U.S. 20004
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Search completed: June 30, 2004, 19:34:04

Job time : 24 secs

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(c) 1993 - 2004 Compugen Ltd.
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protein search, using sw model OM protein

updates/sec Search time 20 Seconds [without alignments] 860.914 Million cell a June 30, 2004, 19:26:48; Run on:

US-10-050-552A-2 903 1 MAALQXSVSSFLMGTLATSC... score: Title: Perfect so Sequence:

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BLOSUM62 Gapop 10.0 , Gapext table: Scoring

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283366 seqs, 96191526 residues Searched:

283366 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

•• Database

PIR 78:*
1: Diri:*
2: Diri:*
3: Diri:*
4: Diri:* 는 0 kg 4i

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

7 0	escription	CRF1 protein - h	erleukin-10 pr	nterleukin-10 pr	erleukin 10 -	ukir	AMP-bi	: abur	il pr	lypro	cal pr	। ਬਸਪੈ	nclec	Ö	cal px	nase (nase (uracil	ical pr	ģ	memb	WD-rep	ikin 10 h	thetical pro	in F09C3.1	slation elo	robable hemag	yclin Gl i	ypothetical pro	protein kinase hom
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ALIGNMENTS

RESULT 2 A38580

interleukin-10 precursor - human
Nighternate names: cytokine synthesis inhibitory factor (CSIF); IL-10
C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 14-Feb-1992 #sequence revision 14-Feb-1992 #text_change 20-Apr-2001
C; Accession: B38580; G01539; Z49110; I37890
R; Vieira, P.; de Waal-Malefyt, R.; Dang, M.N.; Johnson, K.E.; Kastelein, R.; Fiorentin Proc. Natl. Acad. Sci. U.S.A. 88, 1172-1176, 1991
A; Title: Isolation and expression of human cytokine synthesis inhibitory factor cDNA c A; Reference number: A38580; MUID: 91142134; PMID:1847510
A; Accession: A38580
A; Molecule type: mRNA
A; Residues: 1-178 < VIE>

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Query Match
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Matches 40; Conser
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              D:g186271
J.; Baldwin,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PID:9580177
A;Cross-references: GB:M57627; NID:g186270; PIDN:AAA63207.1; FID:g186271
R;Windsor, W.T.: Syco, R: Tasarbopoulos, A.; Zhang, R.; Durkin, J.; Baldw Biochemistry 22, 8807-8815, 1993
A;Title: Disulfide bord assignments and secondary structure analysis of In R;Rahimahal, B.; Mill: A;Rahimahal, B.; Mill: Disulfide bord assignments and secondary structure analysis of In R;Rahimahal, B.; Mill: Singline B.; Mill: Singline B.; Published B.; Published B.; Mill: Singline B.; Status: precombinant protein
R;Raber B.; Ling e.; DNA
A;Residues: Ling e.; DNA
A;Residues: Ling e.; DNA
A;Residues: Ling e.; DNA
A;Residues: Ling e.; Mill: Singline B.; Status, Mill: Singline B.; Singlin
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R; Moore, K.W.; Vieira, P.; Fiorentino, D.F.; Trounstine,
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C;Species: Mus musculus (house mouse)
C;Date: 13-Jul-1990 #sequence_revision 13-Jul-1990
C;Accession: A34853; IS6136
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Pred. No. 0.1;
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ham, T.A.; Moore,
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C; Species: Sus scrofa domestica (domestic pig)
C; Species: Sus scrofa domestica (domestic pig)
C; Date: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_change 26-Aug-1999
C; Accession: I46591
R; Blancho, G.; Gianello, P.; Germana, S.; Baetscher, M.; Sachs, D.H.; LeGuer Proc. Natl. Acad. Sci. U.S.A. 92, 2800-2804, 1995
A; Title: Molecular identification of porcine interleukin 10: regulation of e A; Title: Molecular identification of porcine interleukin 10: regulation of e A; Accession: I46591, MUID: 95224028; PMID: 7708727
A; Accession: I46591
A; Status: preliminary; translated from GB/EMBL/DDBJ
A; Residues: 1-175 <BLA>
A; Cross-references: GB:L20001; NID: g309792; PIDN: AAA74410.1; PID: g309793
C; Genetics:
A; Genetics:
A; Genetics:
A; Genetics:
A; Genetics:
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                                (IL-10)
Science 248, 1230-1234, 1990
A;Title: Homology of cytokine synthesis inhibitory factor (IL-10 A;Reference number: A34853; MUID:90273182; PMID:2161559
A;Accession: A34853
A;Accession: A34853
A;Accession: A34853
A;Molecule type: mRNA
A;Residues: 1-178 «MOO>
A;Cross-references: GB:M37897; NID:g198288; PIDN:AAA39274.1; PID R;Kim, J.M.; Brannan, C.I.; Copeland, N.G.; Jenkins, N.A.; Khan, J. Immunol. 148, 3618-3623, 1992
A;Title: Structure of the mouse IL-10 gene and chromosomal local A;Reference number: I56136; MUID:92268508; PMID:1350294
A;Accession: I56136
A;Accession: I56136
A;Accession: I56136
A;Accession: I56136
A;Generics: DNA
A;Residues: 1-178 «RES>
A;Cross-references: GB:M84340; NID:g198291; PIDN:AAA39275.1; PID C;Genetics:
A;Gene: IL10
A;Introns: 55/3; 75/3; 126/3; 148/3
C;Superfamily: interleukin-10
C;Keywords: cytokine; lymphokine; T-cell
                                                                                                                                                              PIDN:AAA39274.1; PID
Jenkins, N.A.; Khan,
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Pred. No. 0.78;
%; Mismatches
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Pred.
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C; Species: Lactococcus lactis subsp. lactis
C; Species: Lactococcus lactis subsp. lactis
C; Date: 23-Mar-2001 #sequence_revision 23-Mar-2001 #text_change 03-Aug-2001
C; Accession: H86683
R; Bolotin, A.; Wincker, P.; Mauger, S.; Jaillon, O.; Malarme, K.; Weissenbach, J.; Genome Res. 11, 731-753, 2001
A; Title: The complete genome sequence of the lactic acid bacterium Lactococcus lact A; Reference number: A86625; MUID:21235186; PMID:11337471
A; Reference number: A86625; MUID:21235186; PMID:11337471
A; Residues: preliminary
A; Residues: 1-437 <STO>
A; Cross-references: GB:AB005176; PID:g12723351; PIDN:AAK04570.1; GSPDB:GN00146
A; Experimental source: strain IL1403
C; Genetics:
A; Genetics:
                                                                                                                                                                                                                                                 2 AALOKSVSSFIMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLAK
                                                                                 Synechocystis long-chain-fatty-acid-CoA ligase; acetate-CoA
                                                                                                                                                                                                                                                                                                                                                              62 EASLADWNTDVRL--IGE--KLFHGVSMSERCYLMKQVLNFTLEEVLFPQSD-RFQPYMQ
                                                                                                                                                                                                                                                                                                                                                                                                                  -VLDPAVDLRGCEYAO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         embryonic abundant protein EMB24 - white spruce
C;Species: Picea glauca (white spruce)
C;Species: Picea glauca (white spruce)
C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 11-C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 11-C;Accession: T09251
R;Dong, J.Z.; Dunstan, D.I.
R;Dong, J.Z.; Dunstan, D.I.
Submitted to the EMBL Data Library, June 1996
A;Description: Cloning and characterization of 6 novel plant embryog A;Reference number: Z16629
A;Reference number: Z16629
A;Reference number: Z16629
A;Accession: T09251
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Residues: 1-141 < DON>
A;Residues: 1-141 < DON>
A;Genetics:
A;Genetics:
A;Genetics:
A;Genetics:
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A;Genetics:
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                                                                                                                                          Length 601;
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      PID:g15978630;
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                                                                                                                                     DB 2;
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 PIDN: CAC89394.1;
                                                                                                                                     Score 85.5; DB Pred. No. 4.8; 27; Mismatches
                                                                                                                                                                                                                                                                                Score 85; DB 2
Pred, No. 0.93;
5; Mismatches
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Similarity 25.4%;
44; Conservative 27
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iilarity 41.9%;
Conservative 6
 GB:AL590842;
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A;Cross-referenc
C;Genetics:
A;Gene: YP00537
C;Superfamily: S
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C.Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 09-Nov-2001
C.Accession: AG0066
R.Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; Prentice, M.B. deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.; Dougan, G.; il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; Whitehead, S.; Barrell, Nature 413, 523-527, 2001
Nature 413, 523-527, 2001
A; Title: Genome sequence of Yersinia pestis, the causative agent of plague.
A; Reference number: AB0001; MUID:21470413; PMID:11586360
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NATITED THE PROBLES: CYtokine synthesis inhibitory factor
C;Species: Rattus norvegicus (Norway rat)
C;Decession: MNATS; WISSOLI
C;Accession: NUMATS; WISSOLI
R;FENG, L.; Tang, W.W.Y;Chang, J.C.C.; Wilson, C.B.
Sjocher. Biophys. Res. Commun. 192, 452-453, 1992
A;Fitie: Molecular cloning of rat cytokine synthesis inhibitory factor (IL-IC
A;Reference number: Undays
A;Accession: MNATS
A;Accession: MNATS
A;Accession: MNATS
A;Accession: Grad sequence not shown
A;Residues: lu-IC of sequence not shown
A;Residues: Li-IR & FEN.
Biochem. Biophys. Res. Commun. 189, 1-7, 1992
A;Accession: Grad characterization of rat interleukin-IO (IL-IO) cDNA cl
A;Accession: Grass
A;Accession: Grass
A;Accession: Grass
A;Accession: Grass
A;Accession: Grass
A;Accession: S36021
A;Reference number: Grass
A;Accession: S36021
A;Reference number: J36021
A;Reference number: J36021
A;Reference number: J36021
A;Reference number: J36021
A;Residues: Jr-IP F, FBBL: K60675; NID: G2926620; PIDN: CAA43090.1; PID: G296621
C;Superfamily: interleukin-10
C;Superfamily: interleukin-10
C;Superfamily: interleukin-10
C;Superfamily: interleukin-10
C;Superfamily: signal sequence Hstatus predicted ANATS
F;19-178/Product: interleukin-10
F;1-18/Domain: signal sequence Hstatus predicted
F;29,134/Binding site: carbohydrate (Asn) (covalent)
F;29,134/Binding site: carbohydrate (Asn) (covalent)
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LKTLRLRLRRCHQFLPCENKSKAVEEVKSAFSKLQERGVYKAMGBFDI
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C:Species: Yersinia pestis
C:Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_chang
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larity 24.0%; Pred. No. 0.89;
Conservative 33; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12 IMGTLATSCLLLLA-LLVQGGAAAPISSHC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               57 KKDQ--LDNIVLIDSLLQDFKGYLG-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       118 VVPFIARISNRLSTCHIEGDDLH-
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A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-601 < KUI
112
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RESULT 11
G97817
magnesium and cobalt transport protein CorA corA [imported] - Rickettsia concrii (stransport protein CorA corA [imported] - Rickettsia concrii (stransport solute: 30-Sep-2001
C;Species: Rickettsia concrii
C;Date: 30-Sep-2001
A;Ricession: G97817
R;Ogata, H.; Audic, S.; Renesto-Audiffren, P.; Fournier, P.E.; Barbe, V.; Samson, D.; Science 293, 2093-2098, 2001
A;Title: Mechanisms of Evolution in Rickettsia concrii and Rickettsia prowazekii.
A;Reference number: A97700; MUID:21442074; PMID:11557893
A;Accession: G97817
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-319 <KUR>
A;Cross-references: GB:AE006914; PIDN:AAL03481.1; PID:g15620053; GSPDB:GN00173
C;Genetics:
A;Gene: corA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          - FMLAKEASLADNNT-DVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEV
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A; Variety: strain J99
C; Date: 12-Feb-1999 #sequence_revision 12-Feb-1999 #text_charter is 12-Feb-1999 #sequence_revision 12-Feb-1999 #text_charter is 12-Feb-1999 #sequence_revision 12-Feb-1999 #text_charter is 1
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C,Superfamily: polyribonucleotide nucleotidyltransferase
    C;Genetics:
A;Genetic code: SGC3
C;Superfamily: Mycoplasma hypothetical protein MG237
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4.
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Pred. No. 8.2;
1; Mismatches
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Pred. No. 7
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                                                                                                                                                       th 8.8%; Similarity 26.1%; 24; Conservative 18
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Best Local Similarity 19.6%;
Matches 33; Conservative 3
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Best Local S
Matches 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 9
A45389
genome polyprotein - canine distemper virus (strain Onderstepoort)
N'Alternate names: RNA nucleotidyltransferase (RNA-directed); RNA replicase
N'Contains: RNA-directed RNA polymerase (EC 2.7.7.48)
C'Species: canine distemper virus
C'Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 29-May-1998
C'Accession: A45389
R'Sidhu, M.S.; Menonna, J.P.; Cook, S.D.; Dowling, P.C.; Udem, S.A.
Virology 193, 50-65, 1993
A;Title: Canine distemper virus L gene: sequence and comparison with related virusle rate canine distemper virus L gene: sequence and comparison with related vir A;Molecule type: genomic RNA
A;Reference number: A45389; MUID:93174968; PMID:8438585
A;Accession: A45389
A;Accession: A45389
A;Residues: 1-2161 <SID>
C;Genetics:
A;Genetics:
A;Gene: L
C;Superfamily: parainfluenza virus RNA-directed RNA polymerase
C;Keywords: ATP; nucleotidyltransferase; RNA biosynthesis; RNA replication
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C;Species: Mycoplasma genitalium
C;Species: Mycoplasma genitalium
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 0:
C;Accession: B64226
R;Fraser, C.M.; Gocayne, J.D.; White, O.; Adams, M.D.; Clayton, R.
M.; Fuhrmann, J.; Nguyen, D.; Utterback, T.R.; Saudek, D.M.; Phil:
, C.A.; Venter, J.C.
Science 270, 397-403, 1995
A;Title: The minimal gene complement of Mycoplasma genitalium.
A;Reference number: A64200; MUID: 96026346; PMID: 7569993
A;Accession: B64226
A;Accession: B64226
A;Accession: B64226
A;Accession: B0424
A;Residues: 1-294 <TIGR>
A;Residues: 1-294 <TIGR>
A;Experimental source: strain G-37
          <u>ب</u>
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sk, T.R.; Saudek, D.M.; Pr
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H.G.; Glodek, A.; McKenne
.; Bowman, C.; Watthey, L.
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A;Residues: 1-688 <TOM>
A;Cross-references: GB:AE000627; GB:AE000511; NID:g2314373; PIDN:AAD08258.1; PID:g231437
C;Superfamily: polyribonucleotide rucleotidyltransferase
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C; Species: Helicobacter pylori
C; Date: 09-Aug-1997 #sequence_revision 09-Aug-1997 #text_change 27-Oct-2
C; Accession: E64671
R; Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; French, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Feterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glcson, J.D.; Kelley, J.M.; Cotton, M.D.; Weidman, J.M.; Fujii, C.; Bowman, Nature 388, 539-547, 1997
A; Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, A; Title: The complete genome sequence of the gastric pathogen Helicobact A; Reference number: A64520; MUID: 97394467; PMID: 9252185
A; Accession: E64671
A; Status: preliminary; nucleic acid sequence not shown; translation not
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C;Species: Saccharomyces cerevisiae
C;Date: 28-May-1993 #sequence_revision 24-Feb-1995 #text_change
C;Accession: S50608
R;Dietrich, F.S.
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Pred. No. 21;
18; Mismatches
Score 79.5; D
Pred. No. 21;
7; Mismatches
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                                                                 PID:9603343;
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C;Date: 15-Feb-1996 #sequence_revision 01-Mar-1996 #text_change 22-Jun-1999
C;Accession: S59996
R;van den Bergh, F.; Sabina, R.L.
Biochem. J. 312, 401-410, 1995
A;Title: Characterization of human AMP deaminase 2 (AMPD2) gene expression 1
A;Reference number: S5994; MUID:96103174; PMID:8526848
A;Accession: S59996
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Residues: 1-605 <VAN>
A;Residues: 1-605 <VAN>
A;Residues: 1-605 <VAN>
A;Coss-references: EMBL:U16269; NID:g608496; PIDN:AAB06511.1; PID:g608497
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, Octo
C;Genetics:
A;Introns: 175/1; 222/2; 360/3; 404/3
C;Superfamily: AMP deaminase
C;Keywords: alternative splicing; hydrolase
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A; Accession: S50608
A; Molecule type: DNA
A; Residues: 1-1391 < DIE>
A; Residues: 1-1391 < DIE>
A; Cross-references: EMBL: U18839; MID:g603313; PIDN: AAB64660.1;
C; Genetics:
A; Gene: SGD: MUP157
A; Cross-references: SGD: S0000907; MIPS: YER105c
A; Map position: 5R
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Matches 28; Conservative 25; Mismatches
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Pred. No. 22;
28; Mismatches
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Best Local Similarity 25.7%;
Matches 38; Conservative
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/cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

/cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*

/cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

/cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

/cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

/cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

/cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

/cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*

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SUMMARIES	US-10-050-552A-2 US-10-238-965-2 US-09-728-911-15 US-09-870-574-2 US-09-965-528-18 US-09-925-055D-8 US-09-746-375-2 US-10-081-056-354 US-10-232-230-244 US-10-063-745-154 US-10-063-512-154 US-10-063-513-154 US-10-063-513-154	
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          APPLICANT: Colau, Didier
APPLICANT: Colau, Didier
APPLICANT: Renauld, Jean-Christophe
APPLICANT: Dumoutier, Laure
APPLICANT: Dumoutier, Laure
APPLICANT: Polikarpov, Igor
TITLE OF INVENTION: Crystal Structure of Interleukin-22
FILE REFERENCE: LUD-5722 US
CURRENT APPLICATION NUMBER: US/10/050,552A
CURRENT FILING DATE: 2002-04-04
PRIOR APPLICATION NUMBER: 60/317,937
PRIOR FILING DATE: 2001-11-27
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin version 3.1
SEQ ID NO 2
LENGTH: 179
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Pred. No. 9e-92;
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US-10-050-552A-2
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Best Local Similarity
Matches 179; Conser
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; Sequence 2, Application US/10238965
; Publication No. US20040002586A1
; GENERAL INFORMATION:
; APPLICANT: Nagem, Ronaldo A.P.
; APPLICANT: Colau, Didier
; APPLICANT: Colau, Didier
; APPLICANT: Dumoutier, Laure
; APPLICANT: Polikarpov, Igor
; TILE REFERENCE: LUD-5722 US
; TILE REFERENCE: LUD-5722 US
; CURRENT APPLICATION NUMBER: US/10/238,965
; CURRENT FILING DATE: 2001-09-10
; PRIOR FILING DATE: 2001-09-10
; PRIOR FILING DATE: 2001-11-27
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: Patentin version 3.1
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Pred. No. 9e-92;
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APPLICANT: Presnell, Scott R.
APPLICANT: Kindsvogel, Wayne
APPLICANT: Kindsvogel, Wayne
APPLICANT: Chen, Zhi
TITLE OF INVENTIOM: Human Cytokine Receptor
FILE REFERENCE: 99-93
CURRENT APPLICATION NUMBER: US/09/728,911
CURRENT FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: US 60/169,049
PRIOR FILING DATE: 1999-12-03
PRIOR FILING DATE: 2000-09-13
PRIOR FILING DATE: 2000-09-13
PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 36
SOFTWARE: FastSEQ for Windows Version 3.0
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Patent No. US20020012669A1
GENERAL INFORMATION:
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100.0%; Pr
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TYPE: PRT
ORGANISM: Homo sapiens
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APPLICANT: Gurney, Austin L.
APPLICANT: Gurney, Austin L.
APPLICANT: Xie, Ming-Hong
APPLICANT: Xie, Ming-Hong
APPLICANT: Kie, Ming-Hong
APPLICANT: Foster, Jessica S.
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Wood, William I.
TITLE OF INVENTION: INTERLEUKIN-22 POLYPEPTIDES, NUCLEIC ACIDS ENCODING
TITLE OF INVENTION: THE SAME AND METHODS FOR THE TREATMENT OF PANCREATIC DISORDER
TITLE OF INVENTION: THE SAME AND METHODS FOR THE TREATMENT OF PANCREATIC DISORDER
TITLE OF INVENTION NUMBER: US 60/169, 495
CURRENT FILING DATE: 2001-05-30
FRIOR FILING DATE: 2000-05-22
PRIOR PLILNG DATE: 2000-06-24
PRIOR FILING DATE: 2000-06-24
NUMBER: PRIOR FILING DATE: 2000-08-24
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Pred. No. 4.2e-91
; Mismatches
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; Sequence 18, Application US/09965528
; Publication No. US20020187523A1
; GENERAL INFORMATION:
; APPLICANT: INCYTE GENOMICS, INC.
; APPLICANT: TANG, Y. TOM
                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/09870574
Patent No. US20020102723A1
GENERAL INFORMATION:
APPLICANT: Gurney, Austin L.
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il Similarity 98.3%;
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                                                             APPLICANT: BANGHN, Mariah R.
APPLICANT: BAUGHN, Mariah R.
APPLICANT: BAIMZAI, Yalda
APPLICANT: LU, DYUNG Aina M.
APPLICANT: LU, DYUNG Aina M.
APPLICANT: PATTERSON, Chandra
TITLE OF INVENTION: EXTRACELLULAR SIGNALING MOLECULES
FILE REFERENCE: PF-0701 USA
CURRENT APPLICATION NUMBER: US/09/965,528
CURRENT APPLICATION NUMBER: 60/134,949
PRIOR APPLICATION NUMBER: 60/134,949
PRIOR PILING DATE: 1999-07-15
PRIOR APPLICATION NUMBER: 60/146,700
PRIOR APPLICATION NUMBER: 50/146,700
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APPLICANT: Kindsvogel, Wayne R.
APPLICANT: Topouzis, Stavros
TITLE OF INVENTION: SOLUBLE ZCYTOR11 CETCKINE RECEPTORS
FILE REFERENCE: 00-56
CURRENT APPLICATION NUMBER: US/09/925,055D
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: US 60/223,827
PRIOR FILING DATE: 2000-08-08
PRIOR FILING DATE: 2000-12-01
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/ OTHER INFORMATION: Incyte ID No. US20020187523A1
US-09-965-528-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 897; DB 10;
Pred. No. 4.2e-91;
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Pred, No. 4.2e-91
}; Mismatches
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SOFTWARE: FastSEQ for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 8, Application US/09925055D Publication No. US20030157096A1 GENERAL INFORMATION:
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ilarity 98.3%;
Conservative
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                            BURFORD, Neil
BANDMAN, Olga
BAUGHN, Mariah F
AZIMZAI, Yalda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 179
TYPE: PRT
ORGANISM: Homo sapiens
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Best Local Similarity
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ENGTH: 179
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Best Local S
Matches 176
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NAME/KEY:
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    Gaps
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Pred. No. 4.2e-91;
; Mismatches 0;
 Mismatches
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TITLE OF INVENTION: NOVEL CYTOKINE ZCYTO18
FILE REPERENCE: 99-106
CURRENT APPLICATION NUMBER: US/09/746,375
CURRENT FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: US 60/172,105
PRIOR FILING DATE: 1999-12-23
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 44
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 354, Application US/10081056
Publication No. US20040043927A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
                                                                                                                                                                                                                                                      Sequence 2, Application US/09746375
Publication No. US20030170823A1
GENERAL INFORMATION:
APPLICANT: Presnell, Scott R.
APPLICANT: Kindsvogel, Wayne
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Stephan, Jean-Philippe
Watanabe, Colin K.
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Gerber, Hanspeter
Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Hillan, Kenneth J.
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al Similarity 98.3%;
176; Conservative
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Pan, James
Conservative
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US-09-746-375-2
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### APPLICANT: NOOG, WILLIAM. I.

### APPLICANT: ALLIAMS P. NAICH

### APPLICANT: ALLIAMS P. NAICH
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APPLICANT: Baker, Kevin P.
APPLICANT: Besnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Gedard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Austin L.
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Schan, Jean-Philippe F.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530PL60
CURRENT APPLICATION NUMBER: US/10/119,480
FRIOR APPLICATION NUMBER: 60/069113
PRIOR APPLICATION NUMBER: 60/069113
PRIOR APPLICATION NUMBER: 60/063287
PRIOR APPLICATION NUMBER: 60/063287
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-31
PRIOR FILING DATE: 1997-10-31
PRIOR FILING DATE: 1997-10-31
PRIOR FILING DATE: 1997-10-13
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Pred. No. 4.2e-91;
; Mismatches 0;
FRIOR FILING DATE: ZUUL-ULL SONG STO, 574
FRIOR APPLICATION NUMBER: US 09/870,574
FRIOR FILING DATE: 2001-05-30
FRIOR FILING DATE: 2001-05-30
FRIOR FILING DATE: 2001-05-30
FRIOR FILING DATE: 2001-06-01
FRIOR FILING DATE: 2001-06-20
FRIOR FILING DATE: 2001-06-20
FRIOR FILING DATE: 2001-06-20
FRIOR FILING DATE: 2001-06-28
NUMBER OF SEQ ID NOS: 383
SEQ ID NO 354
LENGTH: 179
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Sequence 244, Application US/10219535
Publication No. US20040044179A1
GENERAL INFORMATION:
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FILING DATE: 1997-12-17
APPLICATION NUMBER: 60/078910
FILING DATE: 1998-03-20
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al Similarity 98.3%;
176; Conservative
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FILING DATE: 1998-03-2
APPLICATION NUMBER: 60
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ORGANISM: Homosapiens
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US-10-081-056-354
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RESULT 12
US-10-063-745-154
; Sequence 154, Application US/10063745
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; ORGANISM: Homo S
US-10-232-230-244
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CURRENT APPLICATION NUMBER: US/10/32,230

CURRENT APPLICATION NUMBER: US/119,480

PRIOR APPLICATION NUMBER: 60/059113

PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/062287

PRIOR PILING DATE: 1997-10-17

PRIOR APPLICATION NUMBER: 60/063549

PRIOR APPLICATION NUMBER: 60/063549

PRIOR APPLICATION NUMBER: 60/063549

PRIOR APPLICATION NUMBER: 60/064103
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                                                                                                                                                                       Length 179
                                                 See File Wrapper
                                                                                                                                                                                                    Indels
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                                                                                                                                                                     Score 897; DB 12;
Pred. No. 4.2e-91;
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PRIOR FILING DATE: 1997-10-31
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR FILING DATE: 1998-03-25
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079566
PRIOR FILING DATE: 1998-03-26
PRIOR FILING DATE: 1998-03-26
PRIOR FILING DATE: 1998-3-27
Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 246
SEQ ID NO 244
PRICR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-27
Remaining Prior Application data removed
NUMBER OF SEQ ID MCS: 246
SEQ ID NO 244
LENGTH: 179
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Christopher
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5. US20040044180A1
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Watamabe, Colin L.
Wood, William I.
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Goddard, Audrey
Godowski, Paul J.
Grimaldi, J. Chri
Gurney, Austin L.
Smith, Victoria
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Gerritsen, Mary
                                                                                                                                                                                               Conservative
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                                                                                                      ; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-219-535-244
                                                                                                                                                                                   Similarity
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APPLICANT: Desnoy
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Sequence 244, App
Publication No. 1
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APPLICANT:
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APPLICANT: LAL, Preeti
APPLICANT: BANDMAN, Olga
APPLICANT: BANDMAN, Olga
APPLICANT: BAUGHN, Mariah R.
APPLICANT: BAUGHN, Mariah R.
APPLICANT: AZIMZAI, Yalda
APPLICANT: LU, Dyung Aina M.
APPLICANT: LU, Dyung Aina M.
APPLICANT: PATTERSON, Chandra
TITLE OF INVENTION: EXTRACBLULAR SIGNALING MOLECULES
FILE REFERENCE: PF-0701-1 USA
CURRENT APPLICATION NUMBER: US/09/969,984
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: 60/134,949; 60/144,270; 60/146,700; 60/157,508
NUMBER OF SEQ ID NOS: 55
SOFTWARE: PERL Program
SOFTWARE: PERL Program
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TYPE: PRT

ORGANISM: Homo sapiens

PEATURE:

NAME/KEY: misc feature

OTHER INFORMATION: Incyte ID No. US20040048244A1 5571181CD1

US-09-969-984-18
                                Indels
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Score 897; DB 12;
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Pred. No. 4.2e-91;
3; Mismatches 0;
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APPLICANT: TANG, Y. Tom
APPLICANT: YUE, Henry
APPLICANT: LAL, Preeti
APPLICANT: BURFORD, Neil
APPLICANT: BANDMAN, Olga
APPLICANT: BAUGHN, Mariah R.
APPLICANT: AZIMZAI, Yalda
APPLICANT: LU, Dyung Aina M.
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
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Pred. No. 4.2e-91;
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APPLICANT: Goddard, Audrey
APPLICANT: Goddwski, Paul J.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Watanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRA
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3230R1C1
CURRENT APPLICATION NUMBER: US/12/063,512
CURRENT FILING DATE: 2002-05-01
                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/063,745
CURRENT FILING DATE: 2002-05-09
Prior Application removed - See Palm of SEQ ID NOS: 170
SEQ ID NO 154
LENGTH: 179
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Publication No. US20030018183A1
GENERAL INFORMATION:
APPLICANT: Eaton, Dan L.
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                         Eaton, Dan L.
Filvaroff, Ellen
Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
Grimaldi, Christopher J
Gurney, Austin L.
Watanabe, Colin K.
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Filvaroff, Ellen
Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
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ENGTH: 179
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TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ING THE SAME
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APPLICANT: Goddwski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
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LiCANT: Gerritsen, Mary E.

APPLICANT: Goddard, Audrey

APPLICANT: Godowski, Paul J.

APPLICANT: Grimaldi, Christopher J.

APPLICANT: Gurney, Austin E.

APPLICANT: Watanabe, Colin K.

APPLICANT: Wood, William I.

IITLE OF INVENTION: SECRETED AND TRANSMEMBR.

IITLE OF INVENTION: ACIDS ENCODING THE SAME

FILE REFERENCE: P3230RIC1

CURRENT APPLICATION NUMBER: US/10/063,513

CURRENT APPLICATION TEMOVED - See File W.

NUMBER OF SEQ ID NOS: 170

SEQ ID NO 154

LENGTH: 179

TYPE: PPT
                                                                                                                                                                                                                                                                                                                                                                                                              File Wrapper
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Sequence 154, Application US/10063515
Publication No. US20030018173A1
GENERAL INFORMATION:
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APPLICANT: Filvaroff, Ellen
APPLICANT: Gerritsen, Mary E
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176; Conser
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Publication No. US2
GENERAL INFORMATION
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-10-063-513-154
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Best Local Similarity 98.3%; Pred. No. 4.2e-91;
Matches 176; Conservative 3; Mismatches 0;
CURRENT APPLICATION NUMBER: US/10/063,515
CURRENT FILING DATE: 2002-05-01
Prior Application removed - See File Wrapper NUMBER OF SEQ ID NOS: 170
SEQ ID NO 154
LENGTH: 179
CURRENT AFFLICATION
CURRENT FILING DATE: 20(
Prior Application remove(
NUMBER OF SEQ ID NOS: 17)
SEQ ID NO 154
TENGTH: 179
TYPE: PRT
ORGANISM: Homo Sapien
US-10-053-515-154
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US-10-050-552A-2 903 Title: Perfect score:

1 MAALQKSVSSFLMGTLATSC... Sequence:

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residues 1586107 segs, 282547505 Searched:

1586107 of hits satisfying chosen parameters: Total number

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Minimum Match 0% Maximum Match 100% Listing first 45 summaries

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SUMMARIES

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ALIGNMENTS

TIF; T cell derived inducible factor; interleukin 9; STAT; IL-9; Anti-asthmatic; anti-allergic; cytostatic; inhibitor; antagonist; chromosome 12q15. AAY92879 standard; protein; 179 AA factor. (LUDW-) LUDWIG INST CANCER RES 99WO-US024424. 98US-00178973 99US-00354243 entry) inducible (first WO200024758-A1. Homo sapiens. T cell 18-OCT-1999; 26-OCT-1998; 16-JUL-1999; 04-SEP-2000 04-MAY-2000 AAY92879; Human RESULT 1

٦ .. Renauld Louhed J, Dumoutier L,

N-PSDB; AAA28839, AAA28840. 2000-422495/36

derived inducible factor for cell Wew nucleic acid molecule encoding a T c treating asthma, an allergy or lymphoma.

46pp; English Example 26; Fig 1; This is the human T cell derived inducible factor (TIF). The gene was mapped to chromosome 12q15. The human TIF was identified based on homology to a murine TIF, which was identified by subtraction cloning from a murine lymphoma cell line BW5147 in the presence or absence of interleukin 9 (IL-9). BW5147, can be grown in vitro, without the need to add any cytokines to its culture medium. Many IL-9 activities are mediated by activation of STAT transcription factors. The novel TIFs were expressed in the presence of IL-9, but not in its absence. TIFs induce STAT activation in cells. They can be used, e.g. in the stimulation of regeneration of targeted tissues. Their inhibitors or antagonists can be used to retard, prevent or inhibit differentiation of other tissues. The

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nt of asthma, ridentifying activity in a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides the protein and coding sequences for novel human GIL-19/AE289 protein. The protein shows homology to interleukin-10 (IL-10) and is assumed to be a cytokine. It can be us the regulation of cell proliferation and differentiation, haematopoi immune stimulation or suppression, tissue growth and tumour inhibiti In addition, it also has uses in the treatment of inflammation and i
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TIFs and their coding sequences are useful in the treatment allergies and lymphoma (claimed). They are also useful for icompounds that inhibit or activate T cell induced factor act cell (claimed)
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{interleukin}-10, useful in upregulation of humoral immune
an antiinflammatory agent and as a modulator of immune resj
associated with injury.
                                                                                                                                                                                                                                                                                                                                                        Human; GIL-19/AE289; IL-10; interleukin-10; nutrition; cell proliferation; immune stimulation; immune suppression haematopoiesis regulation; tissue growth; inflammation; ca
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                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
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                                                                     Query Match
Best Local S
Matches 176
                                                                                                                                                                                                                                                                                             AAB36292;
                                                   Sequence
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Human; secreted protein; transmembrane protein; PRO196; PRO444; PRO183; PRO185; PRO210; PRO215; PRO217; PRO242; PRO288; PRO365; PRO1361; PRO1308 PRO1183; PRO1272; PRO1419; PRO4999; PRO7170; PRO248; PRO353; PRO1318; PRO1600; PRO9940; PRO533; PRO301; PRO187; PRO337; PRO1411; PRO4356; PRO246; PRO265; PRO941; PRO10096; PRO6003; PRO6004; PRO350; PRO2630; PRO6309; cell death; genetic disorder; transgenic animal; gene therapy.
                                                                                                  of human polypeptide PRO10096
                                                                                                                                                                                                                                                                                                                                             82. .88 /note= "N-myristoylation site" 97. .10
                                                                                                                                                                                                                                                                                         "N-myristoylation site"
                                                                                                                                                                                                                                                                                                                    site"
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:= "N-glycosylation
                                                                                                                                                                                                                                                                                                                   "N-glycosylation
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                                                                                                                                                                                                                                                                  "signal peptide"
                                                                                                                                                                                                                                         Location/Qualifiers
                         179
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99US-0145070P.
99US-0145698P.
99US-0149396P.
99WO-US020111.
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99WO-US021547.
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2000WO-US004414.
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                       standard; protein;
                                                                           entry)
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/note= "
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20-JUL-1999;
26-JUL-1999;
17-AUG-1999;
01-SEP-1999;
08-SEP-1999;
15-SEP-1999;
15-SEP-1999;
01-DEC-1999;
07-DEC-1999;
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                                                AAB31210;
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RESULT
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179

Length

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897;

Score

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Sequence 179

Query Match

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The present sequence represents a human secreted and transmembrane polypeptide. The specification describes human polypeptides, designated PRO196, PRO136, PRO183, PRO1183, PRO215, PRO217, PRO242, PRO288, PRO365, PRO1361, PRO183, PRO1183, PRO1183, PRO1183, PRO1804, PRO1361, PRO1806, PRO1801, PRO1806, PRO1801, PRO1806, PRO1801, PRO1801,
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Gerritsen ME, Goddard A;
Mather JP, Napier MA, Pan
                                                                                                                                                                                     MA, Par
'illiams
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                                                                                                                                                                                                       3
                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid molecule encoding a PRO polypeptide transmembrane polypeptide is useful for gene therapy and of related polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                Watanabe CK,
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Pred. No. 8.7e-87;
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W, Gerber H,
Kljavin LJ,
                                                                                                                                                                                                 Tumas
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                                                                                                                                                                                                                                                                                                                                                                                                  Claim 12; Fig 64; 244pp; English.
                                                                                                                                                                                                 Stewart TA,
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2000WO-US008439.
2000WO-US013358.
2000WO-US013705.
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986
886
188
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                                                                                                                                        J, Baker KP,
Fong S, Gao ,
Gurney AL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             replace a defective gene
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176; Conservative
                                                                                                (GETH ) GENENTECH INC
                                                                                                                          Ashkena...
Ferrara N, Fo...
Fowski PJ, Gurne,
Roy MA,
                                                                                                                                                                                                                                                          WPI; 2001-050091/06
N-PSDB; AAC87053.
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Ferrara
Godowski PJ,
nni NF, Roy MA
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 20-MAR-2000;
30-MAR-2000;
15-MAY-2000;
17-MAY-2000;
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Best Local S
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Matches
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The patent discloses novel human cytokine, ZCYTO18 protein and its corresponding DNA. ZCYTO18 protein induces proliferation of cells expressing zcytor11, a receptor for ZCYTO18 or induces cytotoxicity in K5626 cells. ZCYTO18 DNA is useful for detecting a genetic abnormality in a patient. ZCYTO18 DNA and its antibodies are useful for detecting cancer and inflammation. ZCYTO18 protein is useful for killing cancer cells. It is useful for increasing platelets in a patient or injured tissue. It is also used in gene therapy. The present sequence is novel human cytokine, ZCYTO18 protein
                                                                                                                                                                                                                                                                 ZCYTO18, useful for treating cancer
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              1, .33
/label= Signal_peptide
34, .179
/label= Mature_ZCYTO18_protein
145. .150
/label= Hydrophilic_region
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Pred. No. 8.7e-87
1; Mismatches
Location/Qualifiers
                                                                                                                                                                                                                                                               Novel human cytokine polypeptide,
                                                                                                                                                                                                                                                                                    4; Page 142; 167pp; English
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01-DEC-2000; 2000US-0250841P
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larity 98.3%;
Conservative
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N-PSDB; AAD09719.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    il Similarity
176; Conser
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Matches 176
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           Peptide
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Human extracellular
  antirheumatic; anti
immunosuppressive;
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Best Local S
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                                                                                                                                                                                                                                                           The present sequence is a human PRO polypeptide (secreted and transmembrane). The PRO protein, and PRO agonists, PRO antagonists or anti-PRO antibodies are useful for preparation of a medicament useful in the treatment of a condition which is responsive to the PRO protein, agonists, antagonists or anti-PRO antibodies. The PRO protein may also be employed as molecular weight markers for protein electrophoresis. The PRO coding sequence has applications in molecular biology, including use as hybridisation probes, and in chromosome and gene mapping
                                                                                                                                                                                                                                                                                                                                                                                                               120
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                                                                                                                                                                     Goddard A
Wood WI;
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Pred. No. 8.7e-87;
Mismatches
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Κ
                                                                                                                                                                     Gerritsen
Watanabe
                                                                                                                                                                                                                                                278pp; English
                            99WO-USG21011.

99WO-USG21090.

99US-0169495P.

2000US-0175481P.

2000WO-USG04341.

2000WO-USG04341.

2000WO-USG04414.

2000WO-USG04414.

2000WO-USG0601.

2000WO-USG08439.

2000US-0191007P.

2000WS-0199397P.

2000WO-USG14042.
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                2000WO-US023328
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larity 98.3%;
Conservative
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including use
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B: AAF92134.
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15-SEP-1999;
07-DEC-1999;
11-JAN-2000;
18-FEB-2000;
18-FEB-2000;
22-FEB-2000;
01-MAR-2000;
03-MAR-2000;
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30-MAR-2000;
30-MAR-2000;
30-MAR-2000;
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              4-AUG-2:0:00;
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gene mapping
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CB-MAR-2001
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                                                                                                                                                                     Eaton DL,
Grimaldi (
                                                                                                                                                                                                                                              Claim 12;
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The invention relates to a human cytokine receptor polypeptide,
designated zcytor16. The zcytor16 polypeptide can be expressed by
standard recombinant methodology and can bind to IL-TIF (undefined). The
standard recombinant methodology and can bind to IL-TIF (undefined). The
zcytor16 protein is useful for: Inhibiting IL-TIF induced proliferation
or differentiation of hematopoietic cell(s) (progenitors); reducing IL-
ITF induced or IL-9 induced inflammation; Heteromeric/multimeric receptor
cresponse in a mammal with inflammation. Heteromeric/multimeric receptor
polypeptides such as soluble zcytor16(CRR2-4 can be used to reduce
progression and symptoms of cancer. Zcytor16 polypeptides can also be
used to detect IL-TIF levels which is indicative of pathological
conditions including inflammatory states (e.g. rheumatory athorities) and
cancer. Antibodies that bind zcytor16 polypeptides and the polypeptides
themselves are useful for the treatment of inflammation, inflammatory
diseases (e.g. infection, asthma, inflammatory bowel disease, rheumatoid
arthritis and atherosclerosis) and autoimmune diseases. The antibodies
and zcytor16 polynucleotides are also useful for detecting cancer. The
present sequence represents the human IL-TIF protein
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Pred. No. 8.7e-87; ; Mismatches 0;
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  antiarthritic;
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larity 98.3%;
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13-SEP-2000;
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Extracellular signaling molecule, EXCS; anti-inflammatory; human, immunosuppressive, cytostatic, neuroprotective, gastrointestinal; virucide, antibacterial; anti-HIV; human immunodeficiency virus; antiinfertility; cerebroprotective, nootropic, antiulcer; antifungal; anticonvulsant; tranquilizer; neuroleptic; vasotropic; gynecological;
                                                                                                            gene therapy
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99US-0144270P.
99US-0146700P.
99US-0157508P.
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                                                                                                          keratolytic; protozoacide;
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Patterson

and polypeptides useful ons and gastrointestinal mmmatory disorders. human extracellular signaling nucleic acids and poly diagnosing, treating and preventing infections and cological, reproductive, and autoimmune/inflammatory neurological,

Claim 1; Page 94; 114pp; English.

The invention provides kuman extracellular signaling molecules (EXCS) and polynucleotides which identify and encode EXCS. EXCS can be expressed by standard recombinant methodology. The amino acid and nucleic acid sequences of EXCS are useful for diagnosing, treating and preventing infections and gastroicrestinal (peptic ulcer, dysphagia, pancreatitis), netrological (e.g. epilepsy, ischemic cerebrovascular disease, stroke), reproductive (infertility, ovulatory defects, endometricois), autoimmune (inflammatory (actinic keratosis, acquired immunodeficiency sprudrome (AIDS), Addison's disease), and cell proliferative disorders including cancers (of the breast, adrenal gland, bone). They may also be used to treat fatal familial insomais, nutritional and metabolic diseases of the cervous system, myopathies, mental disorders (anxiety, schizophrenia, trypanoscas), viral (adenovirus, coronavirus, flavivirus), bacterial (e.g. puramoscus), viral (adenovirus, coronavirus, flavivirus), bacterial (e.g. antagonists, agonists, pharmaceutical compositions, and antibodies may also be used to detect and quantify gene expression increased or decreased expression of EXCS may be correlated with the disease, to determine presence or excess expression of EXCS may bridge with the disease, to determine presence or excess expression of EXCS monitor regulation of EXCS levels dusing therapeutic intervention, to detect the probes, and to detect diseases in microarray, to detect the hybridization probes, and to detect differences in generate hybridization probes, and to detect detect in a finite may and an expression of associated disorders, as targets in microarray. and by ier or affected individuals. Antibodies may lisorders, in monitoring patients being antagonists or inhibitors. Sequences AAB48057 in gene Antibodies may probes, and to detect carrier or affected agonists, antagonists or the EXCS of the invention sequences among normal, carrier or affalso be used in diagnosing disorders, treated with EXCS agonists, antagonist represent also be

Sequence 179 AA;

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                MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                   PJ;
Wood
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Smith V, Stephan
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2000US-0220666P.
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2001WO-US0364208.
2001US-00854208.
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N-PSDB; ABK33657.
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20-MAR-2001;
10-MAY-2001;
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                The invention relates to one hundred and twenty two nucleic acids encoding PRO polypeptides. The sequences of the 122 PRO polypeptides, canced human secreted proteins. The PRO nucleic acids, polypeptides, agonists and antagonists are useful for treating a PRO related disorder. The PRO polypeptides are useful for diagnosing tumours, rectal tumour or liver tumour. The PRO polypeptides are useful for stimulating the proliferation of, or gene expression, in pericyte cells, for stimulating the proliferation or differentiation of chondrocyte cells, for stimulating the release of tumour necrosis factor-alpha from human blood, the proliferation or inhibiting the proliferation of normal human dermal fibroblast cells. The PRO polypeptide may also be used as molecular weight markers and for tissue typing. The PRO nucleic acids have applications in molecular biology, including use as hybridisation probes, and in chromosome and gene mapping. AAUB3592-AAUB3713 represent human PRO
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Pred. No. 8.7e-87;
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B; AAD30645.
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                                                                                          The present sequence is human TIF
                            The invention relates to nucleic acid molecules encoding T cell derived inducible factors (TIFs) also known as interleukin-21 (IL-21). TIF polynucleotides are upregulated by the cytokine, IL-9. IL-TIF or IL-21 molecules are implicated in activation of STAT transcription factors, acute phase proteins and inflammation. The present sequence is human TIP protein. The TIF gene is located on chromosome 12
                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                              Human; secreted protein; transmembrane protein; antirheumatic; antiarthritic; osteopathic; sports-related joint problem; articular cartilage defect; osteoarthritis; rheumatoid arthritis.
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                                                                                                                                                                      Length 179
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                                                                                                                                                                      Score 897; DB 5;
Pred. No. 8.7e-87
}; Mismatches
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 64pp; English
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98US-0083495P.
98US-0087759P.
98US-0088021P.
98US-0088030P.
98US-0088734P.
98US-0088734P.
98US-0088824P.
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98US-0090444P.
98US-0090688P.
98US-0090696F.
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                                                                                                                                                                       99.3%;
98.3%;
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64;
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                                                                                                                                                                                        Similarity
Page
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Gerritsen
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98US-0091628P-
98US-0096012P-
98US-0096757P-
98US-0096757P-
98US-0096949P-
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98US-0099763P-
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98US-0101279P-
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98US-01012733-
98US-0101273P-
98US-0101733P-
98US-0101273P-
98US-0101733P-
98US-01006273-
2000WO-US0233-
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N-PSDB; ABS74454.
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joint problems,
  26-JUN-1998;
10-AUG-1998;
17-AUG-1998;
18-AUG-1998;
26-AUG-1998;
26-AUG-1998;
26-AUG-1998;
26-AUG-1998;
26-AUG-1998;
26-AUG-1998;
26-AUG-1998;
26-AUG-1998;
36-AUG-1998;
36-AUG-1999;
36-AUG-1999;
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PJ. Godowsk Goddard A, Wood WI; A A A

ports-related ful use w ubrane PRO polypeptide us a cell, or for treating rheumatoid arthritis. and transmembrane

sequence of an signal peptide earing as rane PRO The invention relates to an isolated secreted and transmember polypeptide having 80 % sequence identity to a sequence apply ABG95851-ABG95934 or their associated signal peptide, or a extracellular domain of the proteins with their associated

or lacking its associated signal peptide. Also included are the nucleic acids encoding the proteins, vectors, host cells, fusion proteins and useful for detecting a polypeptide designated as A, B, C or D in a sample useful for detecting a polypeptide designated as A, B, C or D in a sample sample with a polypeptide designated as B, F, G, H or D/I polypeptide conjugate in the sample, where the formation of the conjugate is and learnhing the formation of a A/B, B/F, B/G, C/H or D/I polypeptide conjugate in the sample, where the presence of an A, B, C or D polypeptide in the sample, where the presence of an A, B, C or D polypeptide in the sample, where Dolypeptide, B is a PRO1077D polypeptide. It is a PRO1077D polypeptide. The sample conjugate in the sample, where Dolypeptide, B is a PRO1077D polypeptide. The sample conjugate in the sample of the presence of an A, B, C or D polypeptide. H is a PRO1070D polypeptide. The sample congrises a cell suspected of expressing the A, B, C or D polypeptide. The Spide of PRO1070D polypeptide. The sample congrises a cell suspected of expressing the A, B, C or D polypeptide. The sample congrises a cell suspected of expressing a polypeptide designated as A, B, C or D or E, F, G, H or I The bloactive molecule is a toxin, a radiolabel or an antibody. The bloactive molecule sare useful for Inking a biological for a cell expressing a polypeptide designated as A, B, C or D or E, F, G, H, or I The cell is a polypeptide designated as A, B, C or D or E, F, G, H, or I The cell is a toxin, a radiolabel or an antibody. The bloactive molecule is a toxin, a radiolabel or an antibody. The bloactive molecule sains death of the cell. A, B, C, D, B, F, G, H, or I The cell is a polypeptide designated as A, B, C or D or E, F, G, H, or I The cell is a polypeptide designated as A, B, C or D or E, F, G, H, or I The cell is the proteins are useful for activity of a cell expressing a polypeptide designated as A, B, C or D or E, F, G, H, or I The cell is killed. The proteins are useful for activity of a c

Sequence 179 AA;

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                                                MAALOKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                               61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP
                                                                                                61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEBVLFPQSDRFQPYMQEVVP
                                                                                                                                                   FIARISNRLSTCHIRGDDLHIQRNVQKLKDTVKKIGESGBIKAIGELDLLFMSLRNACI
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  Length 179;
                            Indels
Score 897; DB 5; I
Pred. No. 8.7e-87;
I; Mismatches 0;
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99.3%;
                          Conservative
             Similarity
                     176;
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  Match
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Best Local
                       Matches
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ABB84993

ABB84993

standard; protein; 179 16-MAY-2002

Human PRO10096 protein sequence SEQ ID NO:354.

(first entry)

Human; angiogenesis; cardiant; cytostatic; antiangiogenic; hypotensive; vulnerary; antiarteriosclerotic; PRO agonist; PRO antagonist; trauma; gene therapy; cardiovascular disorder; endothelial disorder; cancer; angiogenic disorder; cardiac hypertrophy; atherosclerosis; hypertension; age-related macular degeneration; arterial restenosis; angina; rheumatoid arthritis; myocardial infarction; thrombophlebitis;

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ABEB85003. The PRO proteins given in ABB84817 to ABE885003. The PRO proteins and polynucleotides have cardiant, cytostatic, antiangiogenic, hypotensive, vulnerary and antiarteriosclerotic activities, and can be used in gene therapy. The PRO polynucleotides, proteins, agonists and antagonists are useful for treating or diagnosing a cardiovascular, endothelial or angiogenic disorder in a mammal, e.g. cardiac hypertrophy, trauma, cancer, age-related macular degeneration, atherosclerosis, hypertension, arterial restenosis, rheumatoid arthritis, angina, myocardial infarctions, thrombophlebitis, lymphangitis, tumour angiogenesis (such as breast carcinoma and liver carcinoma) and wound healing. The PRO polynucleotides have applications in molecular biology,
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   carcinoma;
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             mapping
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, Hillan KJ, Marsters
K, Williams PM, Wood W
    east
  breas
gene
  angiogenesis;
some mapping;
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                                                                                                                           2000US-0213637P.
2000US-0220664P.
2000US-0220664P.
2000US-0220664P.
2000US-0222695P.
2000US-0222695P.
2000US-0222695P.
2000US-0222695P.
2000US-0230978P.
2000US-0230978P.
2000US-02643657.
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2000US-02643657.
2000US-02643657.
2000US-02643657.
2000US-02643657.
2000US-0230978P.
2000US-0242922P.
2000US-0242922P.
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2000US-0242922P.
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2000US-0242922P.
2000US-0242922P.
2000US-02429286.
2000US-0266689.
2000US-0266689.
2000US-02866028.
2000US-02866028.
2000US-02866028.
2000US-02866034.
2000US-0287067443.
                                                                                                        2001WO-US01969
             chromosome
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CK,
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Watanabe
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N-PSDB; ABL88248.
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25 - JUL-2000;

25 - JUL-2000;

28 - JUL-2000;

28 - JUL-2000;

28 - JUL-2000;

27 - AUG-2000;

27 - AUG-2000;

28 - AUG-2000;

27 - AUG-2000;

28 - NOV-2000;

28 - NOV-2000;

29 - NOV-2000;

20 - DEC-2000;

20 - DEC-2000;

20 - DEC-2000;

21 - MAR-2001;

22 - MAR-2001;

22 - MAR-2001;

25 - MAY-2001;

25 - MAY-2001;
lymphangitis;
wound healing
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                                                                                                      20-JUN-2001;
                                     Homo sapiens
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01-JUN-2001
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Stephan J
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                                                                                                                     1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                                                             KEASLADNNTDVRLIGEKLFHGVSMSBRCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP
                                                                                                                                                        KEASLADNNTDVFLIGEKLPHGVSMSERCYLMKQVLNFTLEEVLPPQSDRFQPYMQEVVP
                                                                                                    MAALQKSVSFLMGTLATSCLLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
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   nd gene
used in
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  and
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   and in chromosome ar
primers and probes
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                                                              Length
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                                                             Score 897; DB 5; I
Pred. No. 8.7e-87;
; Mismatches 0;
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.e= "Asn is N-myristolated"
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                                                                                                                                                                                                                                                                                                                      protein sequence
  and
  including use as hybridisation probes, an mapping. ABL88259 to ABL88267 represent pexemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                   /label= Signal_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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use as hybridisation
ABL88259 to ABL88267 r
                                                                                                                                                                                                                                                                                                                      Human interleukin 22 (IL-22)
                                                             99.3%;
ilarity 98.3%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  68. .72
/note= "Asn
                                                                                                                                                                                                                                                                                                  (first entry)
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/note= ".
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/note= "
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N-PSDB; ABK11847.
                                                                       Similarity
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                                                                                176;
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Xie M;
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ident for Novel isolated interleukin 22 polypeptide useful for agonists and antagonists that are used for treating chronic pancreatitis, pancreatic carcinoma. Novel isolated agonists and an

ifying IL-22 pancreatitis

Claim 11; Fig 2; 94pp; English

The present invention relates to a new polypeptide having at least 80% identity to a 101 amino acid interleukin (IL)-22 sequence. The invention is useful for detecting IL-22R (IL-22 receptor) or IL-10Rbeta polypeptide in a sample which involves contacting sample with an IL-22 polypeptide and determining the formation of an IL-22R/IL-22 polypeptide conjugate. Preferably, the IL-22 polypeptide or an IL-10Rbeta/IL-22 polypeptide conjugate. Preferably, the IL-22 polypeptide is also useful for linking a bioactive molecule, conjugate. The polypeptide is also useful for linking a bioactive molecule, conjugate or introduced that is bound to the involves contacting the cell with IL-22 polypeptide that is bound to the bioactive molecule and allowing binding of the IL-22 polypeptide with IL-22R or IL-10Rbeta polypeptide thus linking the bioactive molecules to the cell. The molecules of the invention can also be used for modulating biological activity of cell expressing IL-22R or IL-10Rbeta polypeptide. The invention is useful to inhibiting IL-22 induced expression of PAPI (pancreatitis associated to inhibiting IL-22 induced expression of PAPI) or inflamed condition sequence represents for treating pancreatic pulation induced expression of PAF1 (pancreatitic cells. The antibody is also useful fruch as acute or chronic pancreatitis, acinar cell carcinoma or mixed cell pop pancreatic carcinoma and for reducing the activated cell pol of the pancreas in a mammal. The present amino acid sequence human interleukin 22 (IL-22) protein of the invention by pancreatic cells. The ic disorder such as acute including acinar pancreatic carcinoma

Sequence 179 AA;

Ö 120 179 9 9 139 PYITNRIFMLA REOPYMOEVVP Gaps LFMSLRNACI Length 179 KEASLADINTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSD 1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQ MAALQKSVSSFLMGTLATSCLELLALLVQGGAAAPISSHCRLDKSNFQQ **TARISNRLSTCHIEGDDLHIORNVOKLKDTVKKIGESGBIKAIGBLDL** ~ o Score 897; DB 5; Pred. No. 8.7e-87 Mismatches 3, 99.3**%**; 98.3**%**; Conservative Similarity 176; Query Match Best Local S Matches 176 Ç 121 Ġ 121 D C Dp. Ò à ਨੇ 14

standard; protein; (first entry) Human IL-TIF protein #1. 27-DEC-2002 AAE28608; AAE28608 **AAE28608** RESULT

17

Cytokine receptor; Zcytor16; IL-TIF; autoimmune disease; dermatological; inflammatory bowel disease; rheumatoid arthritis; asthma; systemic lupus erythematosus; myasthenia gravis; pancreatitis; diabetes; atherosclerosis; glomerulonephritis; gene therapy; cytostatic; immunosuppressive; nephrotropic; allergy; placental health; abortion; cancer; human.

sapiens Homo

12-SEP-2002

W0200270655-A2

gravis;

2002WO-US006267 04-MAR-2002;

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The invention relates to cytokine receptor designated as mouse Zcytor16 which can bind and antagonise the IL-TIF. The Zcytor16 polypeptide is useful in modulating the immune system by binding Zcytor16 ligand, and thus, preventing the binding of the ligand with endogenous Zcytor16 Library. It is useful for studying human inflammation or immune function, or for treating autoimmune or inflammatory diseases such as inflammatory bowel disease, rheumatoid arthritis, asthma, systemic lupus erythematosus, myasthenia gravis or allergy, atherosclerosis, cancer, diabetes, glomerulonephritis or pancreatitis, or in assessing therapeutic aspects of IL-TIF, chemical therapeutics, anti-IL-TIF antibodies, anti-Zcytor16 antibodies or Zcytor16 soluble receptors. Zcytor16 DNA and the anti-mouse Zcytor16 antibody are useful as probes in detecting gene expression and gene structure, such as in the diagnosis and/or prevention of spontaneous abortions or in monitoring placental health and function. It is also used in gene therapy. The present sequence is human IL-TIF
                                                                                                                                                                      seful for treating autoimmune or inflammatory, bowel disease, rheumatoid arthritis, asthma, diabetes, or in assessing therapeutic aspects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
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                                                                                                                                                                       Zcytor16 polypeptide useful for asee, e.g. inflammatory bowel di
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il Similarity 98.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                   could be expected by the control of the invention related by the collibration related by the collibration related by the collibration related by the collibration of the invention is useful for reducing IL-TIF- or activity. The protein of the invention is useful for reducing IL-TIF- or IL-9 induced inflammation, and inhibiting IL-TIF-induced proliferation. The protein is also useful for suppressing an immune response in a mammal exposed to an antigen or pathogen. Soluble zeytorII receptor or heterodimeric polypeptide is useful for enhancing the in vivo killing of target tissues by directly stimulating a zeytorII receptor-modulated apoptotic pathway. IL-TIF is involved in promoting Thl-type immune responses and antagonists of IL-TIF have beneficial use against diseases involving such immune responses. Soluble zeytorII heterodimers are useful as antagonists in inflammatory and immune disease or conditions such as antagonists in inflammatory and immune disease or conditions such as antagonists, type I diabetes (IDDM), pancreatic cancer, Graves disease, inflammatory bowel disease (IBD), Crohn's disease, colon and intestinal cancer, diverticulosis, autoimmune disease (e.g. IDDW, multiple sclerosis (MS), systemic lupus erythematosus (SiE), myasthaenia gravis, rheumatoid arthritis and IBD), sepsis, asthma, allergy and other atopic diseases, psoriasis and kidney dysfunction. Soluble zeytorII receptor or heterodimeric receptor polypeptides are useful in vivo or in diagnostic cancers in the collibrations and antagonic collibrations and expressing cancers in vivo or in diagnostic capplications to detect IL-TIF expressing cancers in vivo or in diagnostic capplications.
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N-PSDB; ABK10503.
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RESULT 15 ABB95599

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cancer;
disorder;
                                                               Human; angiogenesis; PRO protein; cardiovascularisation; wound;
atherosclerosis; cardiac hypertrophy; gene therapy; endothelial
cardiant; cytostatic; antiangiogenic; hypotensive; vulnerary;
antiarteriosclerotic.
                                               ID NO:
                                                related protein PRO10096
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2000US-0220664P.
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HILLAN K
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32 - MAR - 2001;
10 - MAY - 2001;
25 - MAY - 2001;
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Paoni NF;

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The present invention provides the protein and coding sequences of human PRO proteins. These are useful for treating or diagnosing a cardiovascular, endothelial or angiogenic disorder, including cardiac hypertrophy, trauma, cancer, age-related macular degeneration, atherosclerosis, hypertension, arterial restenosis, rheumatoid arthritis, angina, myocardial infarctions, thrombophlebitis, lymphangitis, tumour angina, myocardial sequence is a PRO protein of the invention
                                                                                                                        PRO polypeptides, [e.g. myocardial a mammal.
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Ye W;
                   Goddard
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               Gerritsen ME, GG
GJ, Marsters SA,
AS PM, Wood WI,
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, Hillan KJ, M:
!K, Williams PM,
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e CK,
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Watanabe
                 Ferrara N,
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ch il Similarity 98.3%; Pred. No. 8.7e-87; 176; Conservative 3; Mismatches 0; Sequence 179 AA;

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